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(54) Title: TIE-2 LIGANDS, METHODS OF MAKING AND USES THEREOF			
(57) Abstract <p>The present invention provides for an isolated nucleic acid molecule encoding a human TIE-2 ligand. In addition, the invention provides for a receptor body which specifically binds a human TIE-2 ligand. The invention also provides an antibody which specifically binds a human TIE-2 ligand. The invention further provides for an antagonist of human TIE-2. The invention also provides for therapeutic compositions as well as a method of blocking blood vessel growth, a method of promoting neovascularization, a method of promoting the growth or differentiation of a cell expressing the TIE-2 receptor, a method of blocking the growth or differentiation of a cell expressing the TIE-2 receptor and a method of attenuating or preventing tumor growth in a human.</p>			

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TIE-2 LIGANDS, METHODS OF MAKING AND USES THEREOF

This Application claims priority of copending International Application PCT US 95/12935, filed October 6, 1995, as well United States Applications U.S. Serial No. 418,595, filed April 6, 1995, the contents of each of which are hereby incorporated by reference. Throughout this application various publications are referenced. The disclosures of these publications in their entireties are hereby incorporated by reference into this application.

INTRODUCTION

The present invention relates generally to the field of genetic engineering and more particularly to genes for receptor tyrosine kinases and their cognate ligands, their insertion into recombinant DNA vectors, and the production of the encoded proteins in recipient strains of microorganisms and recipient eukaryotic cells. More specifically, the present invention is directed to novel ligands, known as the TIE-2 ligands, that bind the TIE-2 receptor, as well as to methods of making and using the TIE-2 ligands. The invention further provides nucleic acid sequences encoding TIE-2 ligands, and methods for the generation of nucleic acids encoding TIE-2 ligands and their gene products. The TIE-2 ligands, as well as nucleic acids encoding them, may be useful in the diagnosis and treatment of certain diseases involving endothelial cells and associated TIE receptors, such as neoplastic diseases involving tumor angiogenesis, wound healing, thromboembolic diseases, atherosclerosis and inflammatory diseases. In addition, the ligands may be used to promote the proliferation and/or differentiation of hematopoietic stem cells.

More generally, biologically active TIE-2 ligands may be used to promote the growth, survival, migration, and/or differentiation and/or stabilization or destabilization of cells expressing the TIE-2 receptor. Biologically active TIE-2 ligand may be used for the in vitro maintenance of TIE-2 receptor expressing cells in culture. Cells and tissues expressing TIE-2 receptor include, for example, cardiac and vascular endothelial cells, lens epithelium and heart epicardium and early hematopoietic cells. Alternatively, such ligand may be used to support cells which are engineered to express TIE-2 receptor. Further, TIE-2 ligands and their cognate receptor may be used in assay systems to identify agonists or antagonists of the TIE-2 receptor.

BACKGROUND OF THE INVENTION

The cellular behavior responsible for the development, maintenance, and repair of differentiated cells and tissues is regulated, in large part, by intercellular signals conveyed via growth factors and similar ligands and their receptors. The receptors are located on the cell surface of responding cells and they bind peptides or polypeptides known as growth factors as well as other hormone-like ligands. The results of this interaction are rapid biochemical changes in the responding cells, as well as a rapid and a long-term readjustment of cellular gene expression. Several receptors associated with various cell surfaces may bind specific growth factors.

The phosphorylation of tyrosine residues in proteins by tyrosine kinases is one of the key modes by which signals are transduced across the plasma membrane. Several currently known protein tyrosine kinase genes encode transmembrane receptors for polypeptide growth factors and hormones such as epidermal growth factor (EGF), insulin, insulin-

like growth factor-I (IGF-I), platelet derived growth factors (PDGF-A and -B), and fibroblast growth factors (FGFs). (Heldin et al., Cell Regulation, 1: 555-566 (1990); Ullrich, et al., Cell, 61: 243-54 (1990)). In each instance, these growth factors exert their action by binding to the extracellular portion of their cognate receptors, which leads to activation of the intrinsic tyrosine kinase present on the cytoplasmic portion of the receptor. Growth factor receptors of endothelial cells are of particular interest due to the possible involvement of growth factors in several important physiological and pathological processes, such as vasculogenesis, angiogenesis, atherosclerosis, and inflammatory diseases. (Folkman, et al. Science, 235: 442-447 (1987)). Also, the receptors of several hematopoietic growth factors are tyrosine kinases; these include c-fms, which is the colony stimulating factor 1 receptor, Sherr, et al., Cell, 41: 665-676 (1985), and c-kit, a primitive hematopoietic growth factor receptor reported in Huang, et al., Cell, 63: 225-33 (1990).

The receptor tyrosine kinases have been divided into evolutionary subfamilies based on the characteristic structure of their ectodomains. (Ullrich, et al. Cell, 61: 243-54 (1990)). Such subfamilies include, EGF receptor-like kinase (subclass I) and insulin receptor-like kinase (subclass II), each of which contains repeated homologous cysteine-rich sequences in their extracellular domains. A single cysteine-rich region is also found in the extracellular domains of the eph-like kinases. Hirai, et al., Science, 238: 1717-1720 (1987); Lindberg, et al. Mol. Cell. Biol., 10: 6316-24 (1990); Lhotak, et al., Mol. Cell. Biol. 11: 2496-2502 (1991). PDGF receptors as well as c-fms and c-kit receptor tyrosine kinases may be grouped into subclass III; while the FGF receptors form subclass IV. Typical for the members of both of these subclasses are extracellular folding units stabilized by intrachain

disulfide bonds. These so-called immunoglobulin (Ig)-like folds are found in the proteins of the immunoglobulin superfamily which contains a wide variety of other cell surface receptors having either cell-bound or soluble ligands. Williams, et al., Ann. Rev. Immunol., 6: 381-405 (1988).

Receptor tyrosine kinases differ in their specificity and affinity. In general, receptor tyrosine kinases are glycoproteins which consist of (1) an extracellular domain capable of binding the specific growth factor(s); (2) a transmembrane domain which usually is an alpha-helical portion of the protein; (3) a juxtamembrane domain where the receptor may be regulated by, e.g., protein phosphorylation; (4) a tyrosine kinase domain which is the enzymatic component of the receptor; and (5) a carboxyterminal tail which in many receptors is involved in recognition and binding of the substrates for the tyrosine kinase.

Processes such as alternative exon splicing and alternative choice of gene promoter or polyadenylation sites have been reported to be capable of producing several distinct polypeptides from the same gene. These polypeptides may or may not contain the various domains listed above. As a consequence, some extracellular domains may be expressed as separate, secreted proteins and some forms of the receptors may lack the tyrosine kinase domain and contain only the extracellular domain inserted in the plasma membrane via the transmembrane domain plus a short carboxyl terminal tail.

A gene encoding an endothelial cell transmembrane tyrosine kinase, originally identified by RT-PCR as an unknown tyrosine kinase-homologous cDNA fragment from human leukemia cells, was described by Partanen, et al., Proc. Natl. Acad. Sci. USA, 87: 8913-8917 (1990). This gene and its encoded protein are called "TIE" which is an

abbreviation for "tyrosine kinase with Ig and EGF homology domains."
Partanen, et al. Mol. Cell. Biol. 12: 1698-1707 (1992).

It has been reported that tie mRNA is present in all human fetal
and mouse embryonic tissues. Upon inspection, tie message has been
5 localized to the cardiac and vascular endothelial cells. Specifically,
tie mRNA has been localized to the endothelia of blood vessels and
endocardium of 9.5 to 18.5 day old mouse embryos. Enhanced tie
expression was shown during neovascularization associated with
developing ovarian follicles and granulation tissue in skin wounds.
10 Korhonen, et al. Blood 80: 2548-2555 (1992). Thus the TIEs has been
suggested to play a role in angiogenesis, which is important for
developing treatments for solid tumors and several other angiogenesis-
dependent diseases such as diabetic retinopathy, psoriasis,
atherosclerosis and arthritis.

15 Two structurally related rat TIE receptor proteins have been
reported to be encoded by distinct genes with related profiles of
expression. One gene, termed tie-1, is the rat homolog of human tie.
Maisonpierre, et al., Oncogene 8: 1631-1637 (1993). The other gene,
tie-2, may be the rat homolog of the murine tek gene, which, like tie,
20 has been reported to be expressed in the mouse exclusively in
endothelial cells and their presumptive progenitors. Dumont, et al.
Oncogene 8: 1293-1301 (1993). The human homolog of tie-2 is
described in Ziegler, U.S. Patent No. 5,447,860 which issued on
September 5, 1995 (wherein it is referred to as "ork"), which is
25 incorporated in its entirety herein.

Both genes were found to be widely expressed in endothelial cells
of embryonic and postnatal tissues. Significant levels of tie-2
transcripts were also present in other embryonic cell populations,

including lens epithelium, heart epicardium and regions of mesenchyme. Maisonpierre, et al., Oncogene 8: 1631-1637 (1993).

The predominant expression of the TIE receptor in vascular endothelia suggests that the TIEs plays a role in the development and maintenance of the vascular system. This could include roles in endothelial cell determination, proliferation, differentiation and cell migration and patterning into vascular elements. Analyses of mouse embryos deficient in TIE-2 illustrate its importance in angiogenesis, particularly for vascular network formation in endothelial cells. Sato, T.N., et al., Nature 376:70-74 (1995). In the mature vascular system, the TIEs could function in endothelial cell survival, maintenance and response to pathogenic influences.

The TIE receptors are also expressed in primitive hematopoietic stem cells, B cells and a subset of megakaryocytic cells, thus suggesting the role of ligands which bind these receptors in early hematopoiesis, in the differentiation and/or proliferation of B cells, and in the megakaryocytic differentiation pathway. Iwama, et al. Biochem. Biophys. Research Communications 195:301-309 (1993); Hashiyama, et al. Blood 87:93-101 (1996), Batard, et al. Blood 87:2212-2220 (1996).

SUMMARY OF THE INVENTION

The present invention provides for a composition comprising a TIE-2 ligand substantially free of other proteins. The invention also provides for an isolated nucleic acid molecule encoding a TIE-2 ligand. The isolated nucleic acid may be DNA, cDNA or RNA. The invention also provides for a vector comprising an isolated nucleic acid molecule encoding a TIE-2 ligand. The invention further provides for a host-vector system for the production in a suitable host cell of a

polypeptide having the biological activity of a TIE-2 ligand. The suitable host cell may be bacterial, yeast, insect or mammalian. The invention also provides for a method of producing a polypeptide having the biological activity of a TIE-2 ligand which comprises growing cells
5 of the host-vector system under conditions permitting production of the polypeptide and recovering the polypeptide so produced.

The invention herein described of an isolated nucleic acid molecule encoding a TIE-2 ligand further provides for the development of the ligand, a fragment or derivative thereof, or another molecule
10 which is a receptor agonist or antagonist, as a therapeutic for the treatment of patients suffering from disorders involving cells, tissues or organs which express the TIE receptor. The present invention also provides for an antibody which specifically binds such a therapeutic molecule. The antibody may be monoclonal or polyclonal. The invention
15 also provides for a method of using such a monoclonal or polyclonal antibody to measure the amount of the therapeutic molecule in a sample taken from a patient for purposes of monitoring the course of therapy.

The present invention also provides for an antibody which specifically binds a TIE-2 ligand. The antibody may be monoclonal or
20 polyclonal. Thus the invention further provides for therapeutic compositions comprising an antibody which specifically binds a TIE-2 ligand in a pharmaceutically acceptable vehicle. The invention also provides for a method of blocking blood vessel growth in a mammal by administering an effective amount of a therapeutic composition
25 comprising an antibody which specifically binds a TIE-2 ligand in a pharmaceutically acceptable vehicle.

The invention further provides for therapeutic compositions comprising a TIE-2 ligand in a pharmaceutically acceptable vehicle.

The invention also provides for a method of promoting neovascularization in a patient by administering an effective amount of a therapeutic composition comprising a TIE-2 ligand in a pharmaceutically acceptable vehicle. In one embodiment, the method
5 may be used to promote wound healing. In another embodiment, the method may be used to treat ischemia. In yet another embodiment, a TIE-2 ligand is used, alone or in combination with other hematopoietic factors, to promote the proliferation or differentiation of hematopoietic stem cells, B cells or megakaryocytic cells.

10 Alternatively, the invention provides that a TIE-2 ligand may be conjugated to a cytotoxic agent and a therapeutic composition prepared therefrom. The invention further provides for a receptorbody which specifically binds a TIE-2 ligand. The invention further provides for therapeutic compositions comprising a receptorbody which specifically
15 binds a TIE-2 ligand in a pharmaceutically acceptable vehicle. The invention also provides for a method of blocking blood vessel growth in a mammal by administering an effective amount of a therapeutic composition comprising a receptorbody which specifically binds a TIE-2 ligand in a pharmaceutically acceptable vehicle.

20 The invention also provides for a TIE-2 receptor antagonist as well as a method of inhibiting TIE-2 ligand biological activity in a mammal comprising administering to the mammal an effective amount of a TIE-2 antagonist. According to the invention, the antagonist may be the TIE-2 ligand 2 as described herein, an antibody or other molecule
25 capable of specifically binding either TIE-2 ligand 1 or TIE-2 receptor (such as for example TIE-2 receptorbody), or ligandbody comprising the fibrinogen-like domain of TIE-2 ligand 1 or ligand 2.

BRIEF DESCRIPTION OF THE FIGURES

FIGURES 1A and 1B - TIE-2 receptorbody (TIE-2 RB) inhibits the development of blood vessels in the embryonic chicken chorioallantoic membrane (CAM). A single piece of resorbable gelatin foam (Gelfoam) soaked with 6 μ g of RB was inserted immediately under the CAM of 1-day chick embryos. After 3 further days of incubation, 4 day old embryos and surrounding CAM were removed and examined. FIGURE 1A: embryos treated with EHK-1 RB (rEHK-1 ecto/hIgG1 Fc) were viable and possessed normally developed blood vessels in their surrounding CAM. FIGURE 1B : all embryos treated with TIE-2 RB (r TIE-2 ecto / h IgG1 Fc) were dead, diminished in size and were almost completely devoid of surrounding blood vessels.

FIGURE 2 - Vector pJFE14.

FIGURE 3 - Restriction map of λ gt10.

FIGURE 4 - Nucleic acid and deduced amino acid (single letter code) sequences of human TIE-2 ligand 1 from clone λ gt10 encoding htie-2 ligand 1.

FIGURE 5 - Nucleic acid and deduced amino acid (single letter code) sequences of human TIE-2 ligand 1 from T98G clone.

FIGURE 6 - Nucleic acid and deduced amino acid (single letter code) sequences of human TIE-2 ligand 2 from clone pBluescript KS encoding human TIE 2 ligand 2.

FIGURE 7 - Western blot showing activation of TIE-2 receptor by TIE-2 ligand 1 (Lane L1) but not by TIE-2 ligand 2 (Lane L2) or control (Mock).

FIGURE 8 - Western blot showing that prior treatment of HAEC cells with excess TIE-2 ligand 2 (Lane 2) antagonizes the subsequent ability of dilute TIE-2 ligand 1 to activate the TIE-2 receptor (TIE2-R) as compared with prior treatment of HAEC cells with MOCK medium (Lane 1).

FIGURE 9- Western blot demonstrating the ability of TL2 to competitively inhibit TL1 activation of the TIE-2 receptor using the human cell hybrid line, EA.hy926.

FIGURE 10- Histogram representation of binding to rat TIE-2 IgG immobilized surface by TIE-2 ligand in C2C12 ras, Rat2 ras, SHEP, and T98G concentrated (10x) conditioned medium. Rat TIE-2 (rTIE2) specific binding is demonstrated by the significant reduction in the binding activity in the presence of 25 µg/ml soluble rat TIE-2 RB as compared to a minor reduction in the presence of soluble trkB RB.

FIGURE 11 - Binding of recombinant human TIE-2 ligand 1 (hTL1) and human TIE-2 ligand 2 (hTL2), in COS cell supernatants, to a human TIE-2 receptorbody (RB) immobilized surface. Human TIE-2-specific binding was determined by incubating the samples with 25 µg/ml of either soluble human TIE-2 RB or trkB RB; significant reduction in the binding activity is observed only for the samples incubated with human TIE-2 RB.

FIGURE 12 - Western blot showing that TIE-2 receptorbody (denoted TIE-2 RB or, as here, TIE2-Fc) blocks the activation of TIE-2 receptors by TIE-2 ligand 1 (TL1) in HUVEC cells, whereas an unrelated receptorbody (TRKB-Fc) does not block this activation.

5

FIGURE 13- Agarose gels showing serial dilutions [undiluted (1) to 10⁻⁴] of the TL1 and TL2 RT-PCR products obtained from E14.5 mouse fetal liver (Lanes 1- total, Lanes 3- stromal enriched, and Lanes 4- c-kit⁺TER119 hematopoietic precursor cells) and E14.5 mouse fetal
10 thymus (Lanes 2- total).

FIGURE 14- Agarose gels showing serial dilutions [undiluted (1) to 10⁻³] of the TL1 and TL2 RT-PCR products obtained from E17.5 mouse fetal thymus cortical stromal cells (Lanes 1- CDR1⁺/A2B5⁻) and medullary
15 stromal cells (Lane CDR1⁻/A2B5⁺).

FIGURE 15- A schematic representation of the hypothesized role of the TIE-2/TIE ligands in angiogenesis. TL1 is represented by (•), TL2 is represented by (*), TIE-2 is represented by (T), VEGF is represented by
20 (□), and flk-1 (a VEGF receptor) is represented by (Y).

FIGURE 16- *In situ* hybridization slides showing the temporal expression pattern of TIE-2, TL1, TL2, and VEGF during angiogenesis associated with follicular development and corpus luteum formation in
25 the ovary of a rat that was treated with pregnant mare serum. Column 1: Early pre-ovulatory follicle; Column 2: pre-ovulatory follicle; Column 3: early corpus luteum; and Column 4: atretic follicle; Row A: bright field; Row B: VEGF; Row C: TL2; Row D: TL1 and Row E: TIE-2 receptor.

FIGURE 17- Diagrammatic representation of the TIE-2 ligands, showing the "coiled coil" and fibrinogen-like domains and the engineering of multimers of the fibrinogen-like domains using antibodies to myc-tags as well as Fc tagging.

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FIGURE 18- Western blot of the covalent multimeric structure of TL1 and TL2 (Panel A) and the interconversion of TL1 and TL2 by the mutation of one cysteine (Panel B).

10 FIGURE 19- A typical curve of TIE-2-IgG binding to immobilized TL1 in a quantitative cell-free binding assay.

FIGURE 20- A typical curve showing TIE-2 ligand 1 ligandbody comprising the fibrinogen-like domain of the ligand bound to the Fc domain of IgG (TL1-fFc) binding to immobilized TIE-2 ectodomain in a
15 quantitative cell-free binding assay.

DETAILED DESCRIPTION OF THE INVENTION

20 As described in greater detail below, applicants have isolated and identified novel ligands that bind the TIE-2 receptor. The TIE-2 ligands of the present invention, which may be purified from nature, or made recombinantly, are referred to herein as TIE-2 ligand 1 (or TL1) and TIE-2 ligand 2 (or TL2). TIE-2 ligand 1, which has an amino acid
25 sequence which is encoded, *inter alia*, by the nucleic acid set forth in Figure 4 or 5, is a TIE-2 receptor agonist. TIE-2 ligand 2, which has an amino acid sequence which is encoded, *inter alia*, by the nucleic acid described in Figure 6, is a TIE-2 receptor antagonist.

The present invention comprises these TIE-2 ligands, as defined by their amino acid sequences, as well as functionally equivalent variants thereof comprising naturally occurring allelic variations, as well as proteins or peptides comprising substitutions, deletions or insertional mutants of the described sequences, which bind the TIE-2 receptors and act as agonists or antagonists thereof. Such variants include those in which amino acid residues are substituted for residues within the sequence resulting in a silent change. For example, one or more amino acid residues within the sequence can be substituted by another amino acid(s) of a similar polarity which acts as a functional equivalent, resulting in a silent alteration. Substitutes for an amino acid within the sequence may be selected from other members of the class to which the amino acid belongs. For example, the class of nonpolar (hydrophobic) amino acids include alanine, leucine, isoleucine, valine, proline, phenylalanine, tryptophan and methionine. The polar neutral amino acids include glycine, serine, threonine, cysteine, tyrosine, asparagine, and glutamine. The positively charged (basic) amino acids include arginine, lysine and histidine. The negatively charged (acidic) amino acids include aspartic acid and glutamic acid.

Also included within the scope of the invention are proteins or fragments or derivatives thereof which exhibit the same or similar biological activity as the TIE-2 ligands described herein, and derivatives which are differentially modified during or after translation, e.g., by glycosylation, proteolytic cleavage, linkage to an antibody molecule or other cellular ligand, etc. Functionally equivalent molecules also include molecules that contain modifications, including N-terminal modifications, which result from expression in a particular recombinant host, such as, for example, N-terminal methylation which occurs in certain bacterial (e.g. E. coli) expression systems. Functional

equivalents also include mutants in which amino acid substitutions are made for cysteine molecules to improve stability of the molecules and to prevent unwanted crosslinking. As used herein, the term "TIE-2 ligand" also include fragments of the TIE-2 ligands which are associated with the binding of the ligands to the TIE-2 receptor. In a preferred embodiment, the TIE-2 ligand comprises the fibrinogen-like domain of TIE-2 ligands 1 and 2 as described herein.

The present invention also encompasses the nucleotide sequence that encodes the proteins described herein as TIE-2 ligands 1 and 2, as well as host cells, including yeast, bacteria, viruses, and mammalian cells, which are genetically engineered to produce the protein, by e.g. transfection, transduction, infection, electroporation, or microinjection of nucleic acid encoding the TIE-2 ligands described herein in a suitable expression vector. The present invention also encompasses introduction of the nucleic acids encoding TIE-2 ligands through gene therapy techniques such as is described, for example, in Finkel and Epstein FASEB J. 9:843-851 (1995); Guzman, et al. PNAS (USA) 91:10732-10736 (1994).

One skilled in the art will also recognize that the present invention encompasses DNA and RNA sequences that hybridize to a deduced TIE-2 ligand encoding sequence, under conditions of moderate stringency, as defined in, for example, Sambrook, et al. Molecular Cloning: A Laboratory Manual, 2 ed. Vol. 1, pp. 101-104, Cold Spring Harbor Laboratory Press (1989). Thus, a nucleic acid molecule contemplated by the invention includes one having a sequence deduced from an amino acid sequence of a TIE-2 ligand prepared as described herein, as well as a molecule having a sequence of nucleic acids that hybridizes to such a nucleic acid sequence, and also a nucleic acid sequence which is degenerate of the above sequences as a result of the

genetic code, but which encodes a ligand that binds the TIE-2 receptor and which has an amino acid sequence and other primary, secondary and tertiary characteristics that are sufficiently duplicative of the ligands described herein so as to confer on the molecule the same biological activity as one of the TIE-2 ligands described herein.

Accordingly, the present invention encompasses an isolated and purified nucleic acid molecule comprising a nucleic acid sequence encoding a human TIE-2 ligand, wherein the nucleic acid sequence is selected from the group consisting of:

- 10 (a) the nucleic acid sequence comprising the coding region of the human TIE-2 ligand as set forth in Figure 4, Figure 5 or Figure 6;
- (b) a nucleic acid sequence that hybridizes under moderately stringent conditions to the nucleic acid sequence of (a) and which encodes a TIE-2 ligand that binds TIE-2 receptor; and
- 15 (c) a nucleic acid sequence that is degenerate as a result of the genetic code to a nucleic acid sequence of (a) or (b), and which encodes a TIE-2 ligand that binds TIE-2 receptor.

The present invention further provides for an isolated and purified human TIE-2 ligand encoded by an isolated nucleic acid molecule of the invention. The invention also provides a vector which comprises an isolated nucleic acid molecule comprising a nucleic acid sequence encoding a human TIE-2 ligand. In one embodiment, the vector is designated as pBluescript KS encoding human TIE 2 ligand 2.

25 Any of the methods known to one skilled in the art for the insertion of DNA fragments into a vector may be used to construct expression vectors encoding TIE-2 ligands using appropriate transcriptional/translational control signals and the protein coding sequences. These methods may include in vitro recombinant DNA and

synthetic techniques and in vivo recombinations (genetic recombination). Expression of a nucleic acid sequence encoding a TIE-2 ligand or peptide fragments thereof may be regulated by a second nucleic acid sequence which is operably linked to the TIE-2 ligand encoding sequence such that the TIE-2 ligand protein or peptide is expressed in a host transformed with the recombinant DNA molecule. For example, expression of a TIE-2 ligand described herein may be controlled by any promoter/enhancer element known in the art. Promoters which may be used to control expression of the ligand include, but are not limited to the long terminal repeat as described in Squinto et al., (Cell 65:1-20 (1991)); the SV40 early promoter region (Bernoist and Chambon, Nature 290:304-310), the CMV promoter, the M-MuLV 5' terminal repeat, the promoter contained in the 3' long terminal repeat of Rous sarcoma virus (Yamamoto, et al., Cell 22:787-797 (1980)), the herpes thymidine kinase promoter (Wagner et al., Proc. Natl. Acad. Sci. U.S.A. 78:144-1445 (1981)), the adenovirus promoter, the regulatory sequences of the metallothionein gene (Brinster et al., Nature 296:39-42 (1982)); prokaryotic expression vectors such as the β -lactamase promoter (Villa-Kamaroff, et al., Proc. Natl. Acad. Sci. U.S.A. 75:3727-3731 (1978)), or the tac promoter (DeBoer, et al., Proc. Natl. Acad. Sci. U.S.A. 80:21-25 (1983)), see also "Useful proteins from recombinant bacteria" in Scientific American, 242:74-94 (1980); promoter elements from yeast or other fungi such as the Gal 4 promoter, the ADH (alcohol dehydrogenase) promoter, PGK (phosphoglycerol kinase) promoter, alkaline phosphatase promoter, and the following animal transcriptional control regions, which exhibit tissue specificity and have been utilized in transgenic animals; elastase I gene control region which is active in pancreatic acinar cells

(Swift et al., Cell 38:639-646 (1984); Ornitz et al., Cold Spring Harbor Symp. Quant. Biol. 50:399-409 (1986); MacDonald, Hepatology 7:425-515 (1987); insulin gene control region which is active in pancreatic beta cells [Hanahan, Nature 315:115-122 (1985)]; immunoglobulin gene
5 control region which is active in lymphoid cells (Grosschedl et al., 1984, Cell 38:647-658; Adames et al., 1985, Nature 318:533-538; Alexander et al., 1987, Mol. Cell. Biol. 7:1436-1444), mouse mammary tumor virus control region which is active in testicular, breast, lymphoid and mast cells (Leder et al., 1986, Cell 45:485-495), albumin
10 gene control region which is active in liver (Pinkert et al., 1987, Genes and Devel. 1:268-276), alpha-fetoprotein gene control region which is active in liver (Krumlauf et al., 1985, Mol. Cell. Biol. 5:1639-1648; Hammer et al., 1987, Science 235:53-58); alpha 1-antitrypsin gene control region which is active in the liver (Kelsey et al., 1987, Genes
15 and Devel. 1:161-171), beta-globin gene control region which is active in myeloid cells (Mogam et al., 1985, Nature 315:338-340; Kollias et al., 1986, Cell 46:89-94); myelin basic protein gene control region which is active in oligodendrocytes in the brain (Readhead et al., 1987, Cell 48:703-712); myosin light chain-2 gene control region which is
20 active in skeletal muscle (Shani, 1985, Nature 314:283-286), and gonadotropic releasing hormone gene control region which is active in the hypothalamus (Mason et al., 1986, Science 234:1372-1378). The invention further encompasses the production of antisense compounds which are capable of specifically hybridizing with a sequence of RNA
25 encoding a TIE-2 ligand to modulate its expression. Ecker, U.S. Patent No. 5,166,195, issued November 24, 1992.

Thus, according to the invention, expression vectors capable of being replicated in a bacterial or eukaryotic host comprising a nucleic acid encoding a TIE-2 ligand as described herein, are used to transfect

a host and thereby direct expression of such nucleic acid to produce the TIE-2 ligand, which may then be recovered in a biologically active form. As used herein, a biologically active form includes a form capable of binding to the TIE-2 receptor and causing a biological response such as a differentiated function or influencing the phenotype of the cell expressing the receptor. Such biologically active forms would, for example, induce phosphorylation of the tyrosine kinase domain of the TIE-2 receptor. Alternatively, the biological activity may be an effect as an antagonist to the TIE-2 receptor, such as is the biological activity of TIE-2 ligand 2. In alternative embodiments, the active form of a TIE-2 ligand is one that can recognize the TIE-2 receptor and thereby act as a targeting agent for the receptor for use in both diagnostics and therapeutics. In accordance with such embodiments, the active form need not confer upon any TIE-2 expressing cell any change in phenotype.

In an alternative embodiment, the biologically active form of the TIE-2 ligand is one in which the ligand is capable of binding the TIE-1 receptor. Preliminary data indicates that TL 2 binds TIE-1 receptor (albeit with low affinity) raising the possibility that it may be able to bind and activate the receptor, or, as in the case of the TIE-2 receptor, bind and act as antagonist.

Expression vectors containing the gene inserts can be identified by four general approaches: (a) DNA-DNA hybridization, (b) presence or absence of "marker" gene functions, (c) expression of inserted sequences and (d) PCR detection. In the first approach, the presence of a foreign gene inserted in an expression vector can be detected by DNA-DNA hybridization using probes comprising sequences that are homologous to an inserted TIE-2 ligand encoding gene. In the second approach, the recombinant vector/host system can be identified and

selected based upon the presence or absence of certain "marker" gene functions (e.g., thymidine kinase activity, resistance to antibiotics, transformation phenotype, occlusion body formation in baculovirus, etc.) caused by the insertion of foreign genes in the vector. For

5 example, if a nucleic acid encoding a TIE-2 ligand is inserted within the marker gene sequence of the vector, recombinants containing the insert can be identified by the absence of the marker gene function. In the third approach, recombinant expression vectors can be identified by assaying the foreign gene product expressed by the recombinant. Such
10 assays can be based, for example, on the physical or functional properties of a TIE-2 ligand gene product, for example, by binding of the ligand to the TIE-2 receptor or portion thereof which may be tagged with, for example, a detectable antibody or portion thereof or by binding to antibodies produced against the TIE-2 ligand protein or a
15 portion thereof. Cells of the present invention may transiently or, preferably, constitutively and permanently express TIE-2 ligands as described herein. In the fourth approach, DNA nucleotide primers can be prepared corresponding to a tie-2 specific DNA sequence. These primers could then be used to PCR a tie-2 gene fragment. (PCR
20 Protocols: A Guide To Methods and Applications, Edited by Michael A. Innis et al., Academic Press (1990)).

The recombinant ligands may be purified by any technique which allows for the subsequent formation of a stable, biologically active protein. Preferably, the ligands are secreted into the culture medium
25 from which they are recovered. Alternatively, the ligand may be recovered from cells either as soluble proteins or as inclusion bodies, from which they may be extracted quantitatively by 8M guanidinium hydrochloride and dialysis in accordance with well known methodology. In order to further purify the ligand, affinity chromatography.

conventional ion exchange chromatography, hydrophobic interaction chromatography, reverse phase chromatography or gel filtration may be used.

In additional embodiments of the invention, as described in greater detail in the Examples, a recombinant TIE-2 ligand encoding gene may be used to inactivate or "knock out" the endogenous gene by homologous recombination, and thereby create a TIE-2 ligand deficient cell, tissue, or animal. For example, and not by way of limitation, the recombinant TIE-2 ligand encoding gene may be engineered to contain an insertional mutation, for example the neo gene, which would inactivate the native TIE-2 ligand encoding gene. Such a construct, under the control of a suitable promoter, may be introduced into a cell, such as an embryonic stem cell, by a technique such as transfection, transduction, or injection. Cells containing the construct may then be selected by G418 resistance. Cells which lack an intact TIE-2 ligand encoding gene may then be identified, e.g. by Southern blotting, PCR detection, Northern blotting or assay of expression. Cells lacking an intact TIE-2 ligand encoding gene may then be fused to early embryo cells to generate transgenic animals deficient in such ligand. Such an animal may be used to define specific in vivo processes, normally dependent upon the ligand.

The present invention also provides for antibodies to the TIE-2 ligands described herein which are useful for detection of the ligands in, for example, diagnostic applications. For preparation of monoclonal antibodies directed toward TIE-2 ligand, any technique which provides for the production of antibody molecules by continuous cell lines in culture may be used. For example, the hybridoma technique originally developed by Kohler and Milstein (1975, Nature 256:495-497), as well as the trioma technique, the human B-cell hybridoma technique (Kozbor

et al., 1983, Immunology Today 4:72), and the EBV-hybridoma technique to produce human monoclonal antibodies (Cole et al., 1985, in "Monoclonal Antibodies and Cancer Therapy," Alan R. Liss, Inc. pp. 77-96) and the like are within the scope of the present invention.

5 The monoclonal antibodies may be human monoclonal antibodies or chimeric human-mouse (or other species) monoclonal antibodies. Human monoclonal antibodies may be made by any of numerous techniques known in the art (e.g., Teng et al., 1983, Proc. Natl. Acad. Sci. U.S.A. 80:7308-7312; Kozbor et al., 1983, Immunology Today 4:72-79; 10 Olsson et al., 1982, Meth. Enzymol. 92:3-16). Chimeric antibody molecules may be prepared containing a mouse antigen-binding domain with human constant regions (Morrison et al., 1984, Proc. Natl. Acad. Sci. U.S.A. 81:6851, Takeda et al., 1985, Nature 314:452).

 Various procedures known in the art may be used for the 15 production of polyclonal antibodies to epitopes of the TIE-2 ligands described herein. For the production of antibody, various host animals, including but not limited to rabbits, mice and rats can be immunized by injection with a TIE-2 ligand, or a fragment or derivative thereof. Various adjuvants may be used to increase the immunological response, 20 depending on the host species, and including but not limited to Freund's (complete and incomplete), mineral gels such as aluminum hydroxide, surface active substances such as lysolecithin, pluronic polyols, polyanions, peptides, oil emulsions, keyhole limpet hemocyanins, dinitrophenol, and potentially useful human adjuvants such as BCG 25 (Bacille Calmette-Guerin) and Corynebacterium parvum.

 A molecular clone of an antibody to a selected TIE-2 ligand epitope can be prepared by known techniques. Recombinant DNA methodology (see e.g., Maniatis et al., 1982, Molecular Cloning, A Laboratory Manual, Cold Spring Harbor Laboratory, Cold Spring Harbor,

New York) may be used to construct nucleic acid sequences which encode a monoclonal antibody molecule, or antigen binding region thereof.

The present invention provides for antibody molecules as well as
5 fragments of such antibody molecules. Antibody fragments which contain the idiotype of the molecule can be generated by known techniques. For example, such fragments include but are not limited to: the $F(ab')_2$ fragment which can be produced by pepsin digestion of the antibody molecule; the Fab' fragments which can be generated by
10 reducing the disulfide bridges of the $F(ab')_2$ fragment, and the Fab fragments which can be generated by treating the antibody molecule with papain and a reducing agent. Antibody molecules may be purified by known techniques, e.g., immunoabsorption or immunoaffinity chromatography, chromatographic methods such as HPLC (high
15 performance liquid chromatography), or a combination thereof.

The present invention further encompasses an immunoassay for measuring the amount of a TIE-2 ligand in a biological sample by
a) contacting the biological sample with at least one antibody which specifically binds the TIE-2 ligand so that the antibody forms a
20 complex with any TIE-2 ligand present in the sample; and
b) measuring the amount of the complex and thereby measuring the amount of the TIE-2 ligand in the biological sample.

The invention further encompasses an assay for measuring the amount of TIE-2 receptor in a biological sample by

25 a) contacting the biological sample with at least one ligand of the invention so that the ligand forms a complex with the TIE-2 receptor; and

- b) measuring the amount of the complex and thereby measuring the amount of the TIE-2 receptor in the biological sample.

The present invention also provides for the utilization of a TIE-2 ligand to support the survival and/or growth and/or migration and/or differentiation of TIE-2 receptor expressing cells. Thus, the ligand may be used as a supplement to support, for example, endothelial cells in culture.

Further, the discovery by applicants of a cognate ligand for the TIE-2 receptor enables the utilization of assay systems useful for the identification of agonists or antagonists of the TIE-2 receptor. Such assay systems would be useful in identifying molecules capable of promoting or inhibiting angiogenesis. For example, in one embodiment, antagonists of the TIE-2 receptor may be identified as test molecules that are capable of interfering with the interaction of the TIE-2 receptor with a biologically active TIE-2 ligand. Such antagonists are identified by their ability to 1) block the binding of a biologically active TIE-2 ligand to the receptor as measured, for example, using BIAcore biosensor technology (BIAcore; Pharmacia Biosensor, Piscataway, NJ); or 2) block the ability of a biologically active TIE-2 ligand to cause a biological response. Such biological responses include, but are not limited to, phosphorylation of the TIE-2 receptor or downstream components of the TIE-2 signal transduction pathway, or survival, growth or differentiation of TIE-2 receptor bearing cells.

In one embodiment, cells engineered to express the TIE-2 receptor may be dependent for growth on the addition of TIE-2 ligand. Such cells provide useful assay systems for identifying additional agonists of the TIE-2 receptor, or antagonists capable of interfering with the activity of TIE-2 ligand on such cells. Alternatively, autocrine cells, engineered to be capable of co-expressing both TIE-2

ligand and receptor, may provide useful systems for assaying potential agonists or antagonists.

Therefore, the present invention provides for introduction of the TIE-2 receptor into cells that do not normally express this receptor, thus allowing these cells to exhibit profound and easily distinguishable responses to a ligand which binds this receptor. The type of response elicited depends on the cell utilized, and not the specific receptor introduced into the cell. Appropriate cell lines can be chosen to yield a response of the greatest utility for assaying, as well as discovering, molecules that can act on tyrosine kinase receptors. The molecules may be any type of molecule, including but not limited to peptide and non-peptide molecules, that will act in systems to be described in a receptor specific manner.

One of the more useful systems to be exploited involves the introduction of the TIE-2 receptor (or a chimeric receptor comprising the extracellular domain of another receptor tyrosine kinase such as, for example, trkC and the intracellular domain of a TIE receptor) into a fibroblast cell line (e.g., NIH3T3 cells) thus such a receptor which does not normally mediate proliferative or other responses can, following introduction into fibroblasts, nonetheless be assayed by a variety of well established methods to quantitate effects of fibroblast growth factors (e.g. thymidine incorporation or other types of proliferation assays; see van Zoelen, 1990, "The Use of Biological Assays For Detection Of Polypeptide Growth Factors" in Progress Factor Research, Vol. 2, pp. 131-152; Zhan and M. Goldfarb, 1986, Mol. Cell. Biol., Vol. 6, pp. 3541-3544). These assays have the added advantage that any preparation can be assayed both on the cell line having the introduced receptor as well as the parental cell line lacking the receptor; only

specific effects on the cell line with the receptor would be judged as being mediated through the introduced receptor. Such cells may be further engineered to express the TIE-2 ligands, thus creating an autocrine system useful for assaying for molecules that act as
5 antagonists/agonists of this interaction. Thus, the present invention provides for host cells comprising nucleic acid encoding TIE-2 ligands and nucleic acid encoding TIE-2 receptor.

The TIE-2 receptor/TIE-2 ligand interaction also provides a useful system for identifying small molecule agonists or antagonists of
10 the TIE-2 receptor. For example, fragments, mutants or derivatives of a TIE-2 ligand may be identified that bind the TIE-2 receptor but do not induce any other biological activity. Alternatively, the characterization of a TIE-2 ligand enables the determination of active portions of the molecule. Further, the identification of a ligand enables
15 the determination of the X-ray crystal structure of the receptor/ligand complex, thus enabling identification of the binding site on the receptor. Knowledge of the binding site will provide useful insight into the rational design of novel agonists and antagonists.

The specific binding of a test molecule to the TIE-2 receptor may
20 be measured in a number of ways. For example, the actual binding of test molecule to cells expressing TIE-2 may be detected or measured, by detecting or measuring (i) test molecule bound to the surface of intact cells; (ii) test molecule cross-linked to TIE-2 protein in cell lysates; or (iii) test molecule bound to TIE-2 in vitro. The specific
25 interaction between test molecule and TIE-2 may be evaluated by using reagents that demonstrate the unique properties of that interaction.

As a specific, nonlimiting example, the methods of the invention may be used as follows. Consider a case in which the TIE-2 ligand in a sample is to be measured. Varying dilutions of the sample (the test
30 molecule), in parallel with a negative control (NC) containing no TIE-2

ligand activity, and a positive control (PC) containing a known amount of a TIE-2 ligand, may be exposed to cells that express TIE-2 in the presence of a detectably labeled TIE-2 ligand (in this example, radioiodinated ligand). The amount of TIE-2 ligand in the test sample
5 may be evaluated by determining the amount of ^{125}I -labeled TIE-2 ligand that binds to the controls and in each of the dilutions, and then comparing the sample values to a standard curve. The more TIE-2 ligand in the sample, the less ^{125}I -ligand that will bind to TIE-2.

The amount of ^{125}I -ligand bound may be determined by measuring
10 the amount of radioactivity per cell, or by cross-linking the TIE-2 ligand to cell surface proteins using DSS, as described in Meakin and Shooter, 1991, Neuron 6:153-163, and detecting the amount of labeled protein in cell extracts using, for example, SDS polyacrylamide gel electrophoresis, which may reveal a labeled protein having a size
15 corresponding to TIE-2 receptor/TIE-2 ligand. The specific test molecule/TIE-2 interaction may further be tested by adding to the assays various dilutions of an unlabeled control ligand that does not bind the TIE-2 receptor and therefore should have no substantial affect on the competition between labeled TIE-2 ligand and test molecule for
20 TIE-2 binding. Alternatively, a molecule known to be able to disrupt TIE-2 receptor/TIE-2 ligand binding, such as, but not limited to, anti-TIE-2 antibody, or TIE-2 receptorbody as described herein, may be expected to interfere with the competition between ^{125}I -TIE-2 ligand and test molecule for TIE-2 receptor binding.

25 Detectably labeled TIE-2 ligand includes, but is not limited to, TIE-2 ligand linked covalently or noncovalently to a radioactive substance, a fluorescent substance, a substance that has enzymatic activity, a substance that may serve as a substrate for an enzyme (enzymes and substrates associated with colorimetrically detectable

reactions are preferred) or to a substance that can be recognized by an antibody molecule that is preferably a detectably labeled antibody molecule.

Alternatively, the specific binding of test molecule to TIE-2 may
5 be measured by evaluating the secondary biological effects of TIE-2
ligand/TIE-2 receptor binding, including, but not limited to, cell growth
and/or differentiation or immediate early gene expression or
phosphorylation of TIE-2. For example, the ability of the test molecule
to induce differentiation can be tested in cells that lack tie-2 and in
10 comparable cells that express tie-2; differentiation in tie-2-
expressing cells but not in comparable cells that lack tie-2 would be
indicative of a specific test molecule/TIE-2 interaction. A similar
analysis could be performed by detecting immediate early gene (e.g. fos
and jun) induction in tie-2-minus and tie-2-plus cells, or by detecting
15 phosphorylation of TIE-2 using standard phosphorylation assays known
in the art. Such analysis might be useful in identifying agonists or
antagonists that do not competitively bind to TIE-2.

Similarly, the present invention provides for a method of
identifying an molecule that has the biological activity of a TIE-2
20 ligand comprising (i) exposing a cell that expresses tie-2 to a test
molecule and (ii) detecting the specific binding of the test molecule to
TIE-2 receptor, in which specific binding to TIE-2 positively correlates
with TIE-2 like activity. Specific binding may be detected by either
assaying for direct binding or the secondary biological effects of
25 binding, as discussed supra. Such a method may be particularly useful
in identifying new members of the TIE ligand family or, in the
pharmaceutical industry, in screening a large array of peptide and non-
peptide molecules (e.g., peptidomimetics) for TIE associated biological
activity. In a preferred, specific, nonlimiting embodiment of the

invention, a large grid of culture wells may be prepared that contain, in alternate rows, PC12 (or fibroblasts, see infra) cells that are either tie-2-minus or engineered to be tie-2-plus. A variety of test molecules may then be added such that each column of the grid, or a portion thereof, contains a different test molecule. Each well could then be scored for the presence or absence of growth and/or differentiation. An extremely large number of test molecules could be screened for such activity in this manner.

In additional embodiments, the invention provides for methods of detecting or measuring TIE ligand-like activity or identifying a molecule as having such activity comprising (i) exposing a test molecule to a TIE-2 receptor protein in vitro under conditions that permit binding to occur and (ii) detecting binding of the test molecule to the TIE-2 receptor protein, in which binding of test molecule to TIE-2 receptor correlates with TIE ligand-like activity. According to such methods, the TIE-2 receptor may or may not be substantially purified, may be affixed to a solid support (e.g. as an affinity column or as an ELISA assay), or may be incorporated into an artificial membrane. Binding of test molecule to TIE-2 receptor may be evaluated by any method known in the art. In preferred embodiments, the binding of test molecule may be detected or measured by evaluating its ability to compete with detectably labeled known TIE-2 ligands for TIE-2 receptor binding.

The present invention also provides for a method of detecting the ability of a test molecule to function as an antagonist of TIE ligand-like activity comprising detecting the ability of the molecule to inhibit an effect of TIE ligand binding to TIE-2 receptor on a cell that expresses the receptor. Such an antagonist may or may not interfere with TIE-2 receptor/TIE-2 ligand binding. Effects of TIE-2 ligand

binding to TIE-2 receptor are preferably biological or biochemical effects, including, but not limited to, cell survival or proliferation, cell transformation, immediate early gene induction, or TIE-2 phosphorylation.

- 5 The invention further provides for both a method of identifying antibodies or other molecules capable of neutralizing the ligand or blocking binding to the receptor, as well as the molecules identified by the method. By way of nonlimiting example, the method may be performed via an assay which is conceptually similar to an ELISA assay.
- 10 For example, TIE-2 receptorbody may be bound to a solid support, such as a plastic multiwell plate. As a control, a known amount of TIE-2 ligand which has been Myc-tagged may then be introduced to the well and any tagged TIE ligand which binds the receptorbody may then be identified by means of a reporter antibody directed against the Myc-tag.
- 15 This assay system may then be used to screen test samples for molecules which are capable of i) binding to the tagged ligand or ii) binding to the receptorbody and thereby blocking binding to the receptorbody by the tagged ligand. For example, a test sample containing a putative molecule of interest together with a known
- 20 amount of tagged ligand may be introduced to the well and the amount of tagged ligand which binds to the receptorbody may be measured. By comparing the amount of bound tagged ligand in the test sample to the amount in the control, samples containing molecules which are capable of blocking ligand binding to the receptor may be identified. The
- 25 molecules of interest thus identified may be isolated using methods well known to one of skill in the art.

Once a blocker of ligand binding is found, one of skill in the art would know to perform secondary assays to determine whether the blocker is binding to the receptor or to the ligand, as well as assays to

determine if the blocker molecule can neutralize the biological activity of the ligand. For example, by using a binding assay which employs BIAcore biosensor technology (or the equivalent), in which either TIE-2 receptorbody or TIE-2 ligand or ligandbody is covalently attached to a solid support (e.g. carboxymethyl dextran on a gold surface), one of skill in the art would be able to determine if the blocker molecule is binding specifically to the ligand, ligandbody or to the receptorbody. To determine if the blocker molecule can neutralize the biological activity of the ligand, one of skill in the art could perform a phosphorylation assay (see Example 5) or alternatively, a functional bioassay, such as a survival assay, by using primary cultures of, for example, endothelial cells. Alternatively, a blocker molecule which binds to the receptorbody could be an agonist and one of skill in the art would know to how to determine this by performing an appropriate assay for identifying additional agonists of the TIE-2 receptor.

In addition, the invention further contemplates compositions wherein the TIE-2 ligand is the receptor binding domains of the TIE-2 ligands described herein. For example, TIE-2 ligand 1 consists of a "coiled coil" domain (beginning at the 5' end and extending to the nucleotide at about position 1160 of Figure 4 and about position 1157 of Figure 5) and a fibrinogen-like domain (which is encoded by the nucleotide sequence of Figure 4 beginning at about position 1161 and about position 1158 of Figure 5). The fibrinogen-like domain of TIE-2 ligand 2 is believed to begin on or around the same amino acid sequence as in ligand 1 (FRDCA) which is encoded by nucleotides beginning around 1197 of Figure 6. Multimerization of the coiled coil domains during production of the ligand hampers purification. As described in Example 19, Applicants have discovered, however, that the fibrinogen-like domain comprises the TIE-2 receptor binding domain. The monomeric

forms of the fibrinogen-like domain do not, however, appear to bind the receptor. Studies utilizing myc-tagged fibrinogen-like domain, which has been "clustered" using anti-myc antibodies, do bind the TIE-2 receptor. [Methods of production of "clustered ligands and ligandbodies are described in Davis, et al. Science 266:816-819 (1994)]. Based on these finding, applicants produced "ligandbodies" which comprise the fibrinogen-like domain of the TIE-2 ligands coupled to the Fc domain of IgG ("fFc's"). These ligandbodies, which form dimers, efficiently bind the TIE-2 receptor. Accordingly, the present invention contemplates the production of TIE-2 ligandbodies which may be used as targeting agents, in diagnostics or in therapeutic applications, such as targeting agents for tumors and/or associated vasculature wherein a TIE-2 antagonist is indicated.

The invention herein further provides for the development of the ligand, a fragment or derivative thereof, or another molecule which is a receptor agonist or antagonist, as a therapeutic for the treatment of patients suffering from disorders involving cells, tissues or organs which express the TIE-2 receptor. Such molecules may be used in a method of treatment of the human or animal body, or in a method of diagnosis.

Because TIE-2 receptor has been identified in association with endothelial cells and, as demonstrated herein, blocking of TIE-2 ligand 1 appears to prevent vascularization, applicants have demonstrated that the TIE-2 ligand 1 will be useful for the induction of vascularization in diseases or disorders where such vascularization is indicated. Such diseases or disorders would include wound healing, ischaemia and diabetes. The ligands may be tested in animal models and used therapeutically as described for other agents, such as vascular endothelial growth factor (VEGF), another endothelial cell-

specific factor that is angiogenic. Ferrara, et al. U.S. Patent No. 5,332,671 issued July 26, 1994. The Ferrara reference, as well as other studies, describe in vitro and in vivo studies that may be used to demonstrate the effect of an angiogenic factor in enhancing blood flow to ischemic myocardium, enhancing wound healing, and in other therapeutic settings wherein neoangiogenesis is desired. [see Sudo, et al. European Patent Application 0 550 296 A2 published July 7, 1993; Banai, et al. Circulation 89:2183-2189 (1994); Unger, et al. Am. J. Physiol. 266:H1588-H1595 (1994); Lazarous, et al. Circulation 91:145-153 (1995)]. According to the invention, the TIE-2 ligands may be used alone or in combination with one or more additional pharmaceutically active compounds such as, for example, VEGF or basic fibroblast growth factor (bFGF), as well as cytokines, neurotrophins, etc.

Conversely, antagonists of the TIE-2 receptor, such as receptorbodies as described herein in Examples 2 and 3, and TIE-2 ligand 2 as described in Example 9, would be useful to prevent or attenuate vascularization, thus preventing or attenuating, for example, tumor growth. These agents may be used alone or in combination with other compositions, such as anti-VEGF antibodies, that have been shown to be useful in treating conditions in which the therapeutic intent is to block angiogenesis. The TIE-2 ligands described herein may also be used in combination with agents, such as cytokine antagonists such as IL-6 antagonists, that are known to block inflammation.

For example, applicants have determined that the TIE-2 ligands are expressed in cells within, or closely associated with, tumors. TIE-2 ligand 2 appears to be tightly associated with tumor endothelial cells. Accordingly, TIE-2 antagonists, such as TIE-2 ligand 2 or TIE-2 ligandbodies may also be useful in preventing or attenuating, for

example, tumor growth. In addition, TIE-2 ligands or ligandbodies may be useful for the delivery of toxins to a receptor bearing cell.

Alternatively, other molecules, such as growth factors, cytokines or nutrients, may be delivered to a TIE-2 receptor bearing cell via TIE-2

5 ligands or ligandbodies. TIE-2 ligands or ligandbodies could also be used as a diagnostic reagent for the TIE-2 receptor, to detect the receptor in vivo or in vitro. Where the TIE-2 receptor is associated with a disease state, TIE-2 ligands or ligandbodies may be useful as diagnostic reagents for detecting the disease by, for example, tissue
10 staining or whole body imaging. Such reagents include radioisotopes, flurochromes, dyes, enzymes and biotin. Such diagnostics or targeting agents may be prepared as described in Alitalo, et al. WO 95/26364 published October 5, 1995 and Burrows, F. and P. Thorpe, PNAS (USA) 90:8996-9000 (1993) which is incorporated herein in its entirety.

15 In other embodiments, the TIE-2 ligands described herein are used as hematopoietic factors. A variety of hematopoietic factors and their receptors are involved in the proliferation and/or differentiation and/or migration of the various cells types contained within blood. Because the TIE-2 receptors are expressed in early hematopoietic
20 cells, the TIE-2 ligands are expected to play a comparable role in the proliferation or differentiation or migration of these cells. Thus, for example, TIE-2 containing compositions may be prepared, assayed, examined in in vitro and in vivo biological systems and used therapeutically as described in any of the following: Sousa, U.S. Patent
25 No. 4,810,643, Lee, et al., Proc. Natl. Acad. Sci. USA 82:4360-4364 (1985) Wong, et al. Science, 228:810-814 (1985); Yokota, et al. Proc. Natl. Acad. Sci (USA) 81:1070 (1984); Bosselman, et al. WO 9105795 published May 2, 1991 entitled "Stem Cell Factor" and Kirkness, et al. WO 95/19985 published July 27, 1995 entitled "Haemopoietic

Maturation Factor". Accordingly, the ligands may be used to diagnose or treat conditions in which normal hematopoiesis is suppressed, including, but not limited to anemia, thrombocytopenia, leukopenia and granulocytopenia. In a preferred embodiment, the TIE-2 ligands may be used to stimulate differentiation of blood cell precursors in situations where a patient has a disease, such as acquired immune deficiency syndrome (AIDS) which has caused a reduction in normal blood cell levels, or in clinical settings in which enhancement of hematopoietic populations is desired, such as in conjunction with bone marrow transplant, or in the treatment of aplasia or myelosuppression caused by radiation, chemical treatment or chemotherapy.

The TIE-2 ligands of the present invention may be used alone, or in combination with another pharmaceutically active agent such as, for example, cytokines, neurotrophins, interleukins, etc. In a preferred embodiment, the ligands may be used in conjunction with any of a number of the above referenced factors which are known to induce stem cell or other hematopoietic precursor proliferation, or factors acting on later cells in the hematopoietic pathway, including, but not limited to, hemopoietic maturation factor, thrombopoietin, stem cell factor, erythropoietin, G-CSF, GM-CSF, etc.

In an alternative embodiment, TIE-2 receptor antagonists, such as TL2, are used to diagnose or treat patients in which the desired result is inhibition of a hematopoietic pathway, such as for the treatment of myeloproliferative or other proliferative disorders of blood forming organs such as thrombocythemias, polycythemias and leukemias. In such embodiments, treatment may comprise use of a therapeutically effective amount of the TIE-2 ligand 2, TIE-2 antibody, TIE-2 receptorbody, a conjugate of a TIE-2 ligand according to claim 27 or 28, or a ligandbody or fFC as described herein.

The present invention also provides for pharmaceutical compositions comprising the TIE-2 ligands or ligandbodies described herein, peptide fragments thereof, or derivatives in a pharmacologically acceptable vehicle. The TIE-2 ligand proteins, peptide fragments, or derivatives may be administered systemically or locally. Any appropriate mode of administration known in the art may be used, including, but not limited to, intravenous, intrathecal, intraarterial, intranasal, oral, subcutaneous, intraperitoneal, or by local injection or surgical implant. Sustained release formulations are also provided for.

The present invention also provides for an antibody which specifically binds such a therapeutic molecule. The antibody may be monoclonal or polyclonal. The invention also provides for a method of using such a monoclonal or polyclonal antibody to measure the amount of the therapeutic molecule in a sample taken from a patient for purposes of monitoring the course of therapy.

The invention further provides for a therapeutic composition comprising a human TIE-2 ligand or ligandbody and a cytotoxic agent conjugated thereto. In one embodiment, the cytotoxic agent may be a radioisotope or toxin.

The invention also provides for an antibody which specifically binds a human TIE-2 ligand. The antibody may be monoclonal or polyclonal.

The invention further provides for a method of purifying a human TIE-2 ligand comprising:

- a) coupling at least one TIE-2 binding substrate to a solid matrix;

- b) incubating the substrate of a) with a cell lysate so that the substrate forms a complex with any human TIE-2 ligand in the cell lysate;
- c) washing the solid matrix; and
- 5 d) eluting the human TIE-2 ligand from the coupled substrate.

The substrate may be any substance that specifically binds the human TIE-2 ligand. In one embodiment, the substrate is selected from the group consisting of anti-TIE-2 ligand antibody, TIE-2
10 receptor and TIE-2 receptorbody. The invention further provides for a receptorbody which specifically binds a human TIE-2 ligand, as well as a therapeutic composition comprising the receptorbody in a pharmaceutically acceptable vehicle, and a method of blocking blood vessel growth in a human comprising administering an effective
15 amount of the therapeutic composition.

The invention also provides for a therapeutic composition comprising a human TIE-2 ligand or ligandbody in a pharmaceutically acceptable vehicle, as well as a method of promoting neovascularization in a patient comprising administering to the
20 patient an effective amount of the therapeutic composition.

In addition, the present invention provides for a method for identifying a cell which expresses TIE-2 receptor which comprises contacting a cell with a detectably labeled TIE-2 ligand or ligandbody, under conditions permitting binding of the detectably labeled ligand
25 to the TIE-2 receptor and determining whether the detectably labeled ligand is bound to the TIE-2 receptor, thereby identifying the cell as one which expresses TIE-2 receptor. The present invention also provides for a therapeutic composition comprising a TIE-2 ligand or

ligandbody and a cytotoxic agent conjugated thereto. The cytotoxic agent may be a radioisotope or toxin.

The invention also provides a method of detecting expression of TIE-2 ligand by a cell which comprises obtaining mRNA from the cell, 5 contacting the mRNA so obtained with a labeled nucleic acid molecule encoding a TIE-2 ligand, under hybridizing conditions, determining the presence of mRNA hybridized to the labeled molecule, and thereby detecting the expression of the TIE-2 ligand in the cell.

The invention further provides a method of detecting expression 10 of a TIE-2 ligand in tissue sections which comprises contacting the tissue sections with a labeled nucleic acid molecule encoding a TIE-2 ligand, under hybridizing conditions, determining the presence of mRNA hybridized to the labelled molecule, and thereby detecting the expression of the TIE-2 ligand in tissue sections.

15

EXAMPLE 1 - IDENTIFICATION OF THE ABAE CELL LINE AS
REPORTER CELLS FOR THE TIE-2 RECEPTOR

Adult BAE cells are registered in the European Cell Culture 20 Repository, under ECACC#92010601. (See PNAS 75:2621 (1978)). Northern (RNA) analyses revealed moderate levels of tie-2 transcripts in the ABAE (Adult Bovine Arterial Endothelial) cell line, consistent with in situ hybridization results that demonstrated almost exclusive localization of tie-2 RNAs to vascular endothelial cells. We therefore 25 examined ABAE cell lysates for the presence of TIE-2 protein, as well as the extent to which this TIE-2 protein is tyrosine-phosphorylated under normal versus serum-deprived growth conditions. ABAE cell lysates were harvested and subjected to immunoprecipitation, followed by Western blot analyses of immunoprecipitated proteins with TIE-2

specific and phosphotyrosine-specific antisera. Omission or inclusion of TIE-2 peptides as specific blocking molecules during TIE-2 immunoprecipitation allowed unambiguous identification of TIE-2 as a moderately detectable protein of ~150 kD whose steady-state
5 phosphotyrosine levels diminish to near undetectable levels by prior serum-starvation of the cells.

Culture of ABAE cells and harvest of cell lysates was done as follows. Low-passage-number ABAE cells were plated as a monolayer at a density of 2×10^6 cells/150mm plastic petri plate (Falcon) and
10 cultured in Dulbecco's modified Eagle's medium (DMEM) containing 10% bovine calf serum (10 % BCS), 2 mM L-glutamine (Q) and 1% each of penicillin and streptomycin (P-S) in an atmosphere of 5% CO₂. Prior to harvest of cell lysates, cells were serum-starved for 24 hours in DMEM/Q/P-S, followed by aspiration of the medium and rinsing of the
15 plates with ice-cold phosphate buffered saline (PBS) supplemented with sodium orthovanadate, sodium fluoride and sodium benzamidine. Cells were lysed in a small volume of this rinse buffer that had been supplemented with 1% NP40 detergent and the protease inhibitors PMSF and aprotinin. Insoluble debris was removed from the cell lysates by
20 centrifugation at 14,000 xG for 10 minutes, at 4°C and the supernatants were subjected to immunoprecipitation with antisera specific for TIE-2 receptor, with or without the presence of blocking peptides added to ~20 µg/ml lysate. Immunoprecipitated proteins were resolved by PAGE (7.5% Laemmli gel), and then electro-transferred to
25 PVDF membrane and incubated either with various TIE-2- or phosphotyrosine-specific antisera. TIE-2 protein was visualized by incubation of the membrane with HRP-linked secondary antisera followed by treatment with ECL reagent (Amersham).

**EXAMPLE 2 - CLONING AND EXPRESSION OF TIE-2 RECEPTOR BODY FOR
AFFINITY-BASED STUDY OF TIE-2 LIGAND
INTERACTIONS**

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An expression construct was created that would yield a secreted protein consisting of the entire extracellular portion of the rat TIE-2 receptor fused to the human immunoglobulin gamma-1 constant region (IgG1 Fc). This fusion protein is called a TIE-2 "receptorbody" (RB), and would be normally expected to exist as a dimer in solution based on formation of disulfide linkages between individual IgG1 Fc tails. The Fc portion of the TIE-2 RB was prepared as follows. A DNA fragment encoding the Fc portion of human IgG1 that spans from the hinge region to the carboxy-terminus of the protein, was amplified from human placental cDNA by PCR with oligonucleotides corresponding to the published sequence of human IgG1; the resulting DNA fragment was cloned in a plasmid vector. Appropriate DNA restriction fragments from a plasmid encoding the full-length TIE-2 receptor and from the human IgG1 Fc plasmid were ligated on either side of a short PCR-derived fragment that was designed so as to fuse, in-frame, the TIE-2 and human IgG1 Fc protein-coding sequences. Thus, the resulting TIE-2 ectodomain-Fc fusion protein precisely substituted the IgG1 Fc in place of the region spanning the TIE-2 transmembrane and cytoplasmic domains. An alternative method of preparing RBs is described in Goodwin, et. al. Cell 73:447-456 (1993).

Milligram quantities of TIE-2 RB were obtained by cloning the TIE-2 RB DNA fragment into the pVL1393 baculovirus vector and subsequently infecting the Spodoptera frugiperda SF-21AE insect cell line. Alternatively, the cell line SF-9 (ATCC Accession No. CRL-1711)

or the cell line BTI-TN-5b1-4 may be used. DNA encoding the TIE-2 RB was cloned as an Eco RI-NotI fragment into the baculovirus transfer plasmid pVL1393. Plasmid DNA purified by cesium chloride density gradient centrifugation was recombined into viral DNA by mixing 3 μ g of plasmid DNA with 0.5 μ g of Baculo-Gold DNA (Pharminigen), followed by introduction into liposomes using 30 μ g Lipofectin (GIBCO-BRL). DNA-liposome mixtures were added to SF-21AE cells (2×10^6 cells/60mm dish) in TMN-FH medium (Modified Grace's Insect Cell Medium (GIBCO-BRL) for 5 hours at 27°C, followed by incubation at 27°C for 5 days in TMN-FH medium supplemented with 5% fetal calf serum. Tissue culture medium was harvested for plaque purification of recombinant viruses, which was carried out using methods previously described (O'Reilly, D.R., L.K. Miller, and V.A. Luckow, Baculovirus Expression Vectors - A Laboratory Manual, 1992, New York: W.H. Freeman) except that the agarose overlay contained 125 μ g/mL X-gal (5-bromo-4-chloro-3-indolyl- β -D-galactopyranoside; GIBCO-BRL). After 5 days of incubation at 27°C, non-recombinant plaques were scored by positive chromogenic reaction to the X-gal substrate, and their positions marked. Recombinant plaques were then visualized by addition of a second overlay containing 100 μ g/mL MTT (3-[4,5-dimethylthiazol-2-yl]2,5-diphenyltetrazolium bromide; Sigma). Putative recombinant virus plaques were picked by plug aspiration, and purified by multiple rounds of plaque isolation to assure homogeneity. Virus stocks were generated by serial, low-multiplicity passage of plaque-purified virus. Low passage stocks of one virus clone (vTIE-2 receptorbody) were produced.

SF-21AE cells were cultured in serum free medium (SF-900 II, Gibco BRL) containing 1X antibiotic/antimycotic solution (Gibco BRL)

and 25 mg/L Gentamycin (Gibco BRL). Pluronic F-68 was added as a surfactant to a final concentration of 1g/L. Cultures (4L) were raised in a bioreactor (Artisan Cell Station System) for at least three days prior to infection. Cells were grown at 27°C, with gassing to 50 %
5 dissolved oxygen, at a gas flow rate of 80 mL/min (aeration at a sparge ring). Agitation was by means of a marine impeller at a rate of 100 rpm. Cells were harvested in mid-logarithmic growth phase ($\sim 2 \times 10^6$ cells/mL), concentrated by centrifugation, and infected with 5 plaque forming units of vTIE-2 receptorbody per cell. Cells and inoculum were
10 brought to 400mL with fresh medium, and virus was adsorbed for 2 hours at 27°C in a spinner flask. The culture was then resuspended in a final volume of 8L with fresh serum-free medium, and the cells incubated in the bioreactor using the previously described conditions.

Culture medium from vTIE-2 receptorbody-infected SF21AE cells
15 were collected by centrifugation (500x g, 10 minutes) at 72 hours post-infection. Cell supernatants were brought to pH 8 with NaOH. EDTA was added to a final concentration of 10 mM and the supernatant pH was readjusted to 8. Supernatants were filtered (0.45 μ m, Millipore) and loaded on a protein A column (protein A sepharose 4 fast
20 flow or HiTrap protein A, both from Pharmacia). The column was washed with PBS containing 0.5 M NaCl until the absorbance at 280 nm decreased to baseline. The column was washed in PBS and eluted with 0.5 M acetic acid. Column fractions were immediately neutralized by eluting into tubes containing 1 M Tris pH 9. The peak fractions
25 containing the TIE-2 receptorbody were pooled and dialyzed versus PBS.

EXAMPLE 3 - **DEMONSTRATION THAT TIE-2 HAS A CRITICAL**
ROLE IN DEVELOPMENT OF THE VASCULATURE

Insight into the function of TIE-2 was gained by introduction of "excess" soluble TIE-2 receptorbody (TIE-2 RB) into a developing system. The potential ability of TIE-2 RB to bind, and thereby neutralize, available TIE-2 ligand could result in an observable
5 disruption of normal vascular development and characterization of the ligand. To examine whether TIE-2 RB could be used to disrupt vascular development in early chick embryos, small pieces of a biologically resorbable foam were soaked with TIE-2 RB and inserted immediately beneath the chorioallantoic membrane at positions just lateral to the
10 primitive embryo.

Early chicken embryos develop atop the yolk from a small disk of cells that is covered by the chorioallantoic membrane (CAM). The endothelial cells that will come to line the vasculature in the embryo arise from both extra- and intra-embryonic cell sources. Extra-
15 embryonically-derived endothelial cells, which provide the major source of endothelial cells in the embryo, originate from accretions of mesenchyme that are situated laterally around the embryo-proper, just underneath the CAM. As these mesenchyme cells mature, they give rise to a common progenitor of both the endothelial and hematopoietic cell
20 lineages, termed the hemangioblast. In turn, the hemangioblast gives rise to a mixed population of angioblasts (the endothelial cell progenitor) and hematoblasts (the pluripotential hematopoietic precursor). Formation of rudiments of the circulatory system begins when endothelial cell progeny segregate to form a one-cell-thick
25 vesicle that surrounds the primitive blood cells. Proliferation and migration of these cellular components eventually produces a vast network of blood-filled microvessels under the CAM that will ultimately invade the embryo to join with limited, intra-embryonically-derived vascular elements.

Newly fertilized chicken eggs obtained from Spafas, Inc. (Boston, MA) were incubated at 99.5°F, 55 % relative humidity. At about 24 hrs. of development, the egg shell was wiped down with 70% ethanol and a dentist's drill was used to make a 1.5 cm. hole in the blunt apex of each egg. The shell membrane was removed to reveal an air space directly above the embryo. Small rectangular pieces of sterile Gelfoam (Upjohn) were cut with a scalpel and soaked in equal concentrations of either TIE-2- or EHK-1 receptorbody. EHK-1 receptorbody was made as set forth in Example 2 using the EHK-1 extracellular domain instead of the TIE-2 extracellular domain (Maisonpierre et al., Oncogene 8:3277-3288 (1993). Each Gelfoam piece absorbed approximately 6 µg of protein in 30 µl. Sterile watchmakers forceps were used to make a small tear in the CAM at a position several millimeters lateral to the primitive embryo. The majority of the piece of RB-soaked Gelfoam was inserted under the CAM and the egg shell was sealed over with a piece of adhesive tape. Other similarly-staged eggs were treated in parallel with RB of the unrelated, neuronally expressed receptor tyrosine kinase, EHK-1 (Maisonpierre et al., Oncogene 8:3277-3288 (1993). Development was allowed to proceed for 4 days and then the embryos were examined by visual inspection. Embryos were removed by carefully breaking the shells in dishes of warmed PBS and carefully cutting away the embryo with surrounding CAM. Of 12 eggs treated with each RB, 6 TIE-2 RB and 5 EHK-1 RB treated embryos had developed beyond the stage observed at the start of the experiment. A dramatic difference was seen between these developed embryos, as shown in Figures 1A and 1B. Those treated with EHK-1 RB appeared to have developed relatively normally. Four out of five EHK-1 embryos were viable as judged by the presence of a beating heart. Furthermore,

the extra-embryonic vasculature, which is visually obvious due to the presence of red blood cells, was profuse and extended several centimeters laterally under the CAM. By contrast, those treated with TIE-2 RB were severely stunted, ranging from 2-5 mm. in diameter, as compared with more than 10 mm in diameter for the EHK-1 RB embryos. All of the TIE-2 RB treated embryos were dead and their CAMs were devoid of blood vessels. The ability of TIE-2 RB to block vascular development in the chicken demonstrates that TIE-2 ligand is necessary for development of the vasculature.

**EXAMPLE 4 - IDENTIFICATION OF A TIE-2-SPECIFIC BINDING
ACTIVITY IN CONDITIONED MEDIUM FROM THE *ras*
ONCOGENE-TRANSFORMED C2C12 MOUSE MYOBLAST
CELL LINE**

Screening of ten-fold-concentrated cell-conditioned media (10X CCM) from various cell lines for the presence of soluble, TIE-2-specific binding activity (BIAcore; Pharmacia Biosensor, Piscataway, NJ) revealed binding activity in serum-free medium from oncogenic-*ras*-transformed C2C12 cells (C2C12-*ras*), RAT 2-*ras* (which is a *ras* transformed fibroblast cell line), human glioblastoma T98G and the human neuroblastoma cell line known as SHEP-1.

The C2C12-*ras* 10X CCM originated from a stably transfected line of C2C12 myoblasts that was oncogenically transformed by transfection with the T-24 mutant of H-*ras* by standard calcium phosphate-based methods. An SV40 based neomycin-resistance expression plasmid was physically linked with the *ras* expression plasmid in order to permit selection of transfected clones. Resulting G418-resistant *ras*-C2C12 cells were routinely maintained as a monolayer on plastic dishes in DMEM/glutamine/penicillin-

streptomycin supplemented with 10 % fetal calf serum (FCS). Serum-free C2C12-~~ras~~ 10X CCM was made by plating the cells at 60% confluence in a serum free defined media for 12 hours. [Zhan and Goldfarb, Mol. Cell. Biol. 6: 3541-3544 (1986)]; Zhan, et al. Oncogene 1: 369-376 (1987)]. The medium was discarded and replaced with fresh DMEM/Q/P-S for 24 hours. This medium was harvested and cells were re-fed fresh DMEM/Q/P-S, which was also harvested after a further 24 hours. These CCM were supplemented with the protease inhibitors PMSF (1mM) and aprotinin (10µg/ml), and ten-fold concentrated on sterile size-exclusion membranes (Amicon). TIE-2-binding activity could be neutralized by incubation of the medium with an excess of TIE-2 RB, but not by incubation with EHK-1 RB, prior to BIAcore analysis.

Binding activity of the 10x CCM was measured using biosensor technology (BIAcore; Pharmacia Biosensor, Piscataway, NJ) which monitors biomolecular interactions in real-time via surface plasmon resonance. Purified TIE-2 RB was covalently coupled through primary amines to the carboxymethyl dextran layer of a CM5 research grade sensor chip (Pharmacia Biosensor; Piscataway, NJ). The sensor chip surface was activated using a mixture of N-hydroxysuccinimide (NHS) and N-ethyl-N'-(3-dimethylaminopropyl)carbodiimide (EDC), followed by immobilization of TIE-2 RB (25 µg/mL, pH 4.5) and deactivation of unreacted sites with 1.0 M ethanolamine (pH 8.5). A negative control surface of the EHK-1 receptorbody was prepared in a similar manner.

The running buffer used in the system was HBS (10 mM Hepes, 3.4 mM EDTA, 150 mM NaCl, 0.005% P20 surfactant, pH 7.4). The 10x CCM samples were centrifuged for 15 min at 4° C and further clarified using a sterile, low protein-binding 0.45 µm filter (Millipore; Bedford, MA).

Dextran (2mg/ml) and P20 surfactant (0.005%) were added to each CCM sample. Aliquots of 40 μ L were injected across the immobilized surface (either TIE-2 or EHK-1) at a flow rate of 5 μ L/min and the receptor binding was monitored for 8 min. The binding activity (resonance units, RU) was measured as the difference between a baseline value determined 30 s prior to the sample injection and a measurement taken at 30 s post-injection. Regeneration of the surface was accomplished with one 12- μ L pulse of 3 M $MgCl_2$.

The instrument noise level is 20 RU; therefore, any binding activity with a signal above 20 RU may be interpreted as a real interaction with the receptor. For C2C12-ras conditioned media, the binding activities were in the range 60-90 RU for the TIE-2 RB immobilized surface. For the same samples assayed on a EHK-1 RB immobilized surface, the measured activities were less than 35 RU.

Specific binding to the TIE-2 receptorbody was evaluated by incubating the samples with an excess of either soluble TIE-2 or EHK-1 RB prior to assaying the binding activity. The addition of soluble EHK-1 RB had no effect on the TIE-2 binding activity of any of the samples, while in the presence of soluble TIE-2 binding to the surface is two-thirds less than that measured in the absence of TIE-2. A repeat assay using >50x concentrated C2C12-ras CCM resulted in a four-fold enhancement over background of the TIE-2 specific binding signal.

EXAMPLE 5 - C2C12-ras CCM CONTAINS AN ACTIVITY THAT INDUCES TYROSINE PHOSPHORYLATION OF TIE-2 RECEPTOR

C2C12-ras 10X CCM was examined for its ability to induce tyrosine phosphorylation of TIE-2 in ABAE cells. Serum-starved ABAE cells were briefly incubated with C2C12-ras CCM, lysed and subjected

to immunoprecipitation and Western analyses as described above.

Stimulation of serum-starved ABAE cells with serum-free C2C12-ras 10X CCM was done as follows. The medium of ABAE cells starved as described above was removed and replaced with either defined medium
5 or 10X CCM that had been pre-warmed to 37°C. After 10 minutes, the media were removed and the cells were twice rinsed on ice with an excess of chilled PBS supplemented with orthovanadate/NaF/benzamidine. Cell lysis and TIE-2-specific immunoprecipitation was done as described above.

10 ABAE cells incubated for 10 minutes with defined medium showed no induction of TIE-2 tyrosine phosphorylation, whereas incubation with C2C12-ras CCM stimulated at least a 100 X increase in TIE-2 phosphorylation. This activity was almost totally depleted by pre-incubation of the C2C12-ras 10X CCM for 90 minutes at room
15 temperature with 13 µg of TIE-2 RB coupled to protein G-Sepharose beads. Medium incubated with protein G Sepharose alone was not depleted of this phosphorylating activity.

20 EXAMPLE 6 - EXPRESSION CLONING OF TIE-2 LIGAND

COS-7 cells were cultured in Dulbecco's modified Eagle's medium (DMEM) containing 10% fetal bovine serum (FBS), 1% each of penicillin and streptomycin (P/S) and 2 mM glutamine in an atmosphere of 5% CO₂. The mouse myoblast C2C12 ras cell line was cultured in Eagle's
25 minimal essential medium (EMEM) with 10% FBS, (P/S) and 2 mM glutamine. Full length mouse TIE-2 ligand cDNA clones were obtained by screening a C2C12 ras cDNA library in the pJFE14 vector expressed in COS cells. This vector, as shown in Figure 2, is a modified version of

the vector pSR_α (Takebe, et al. 1988, Mol. Cell. Biol. 8:466-472). The library was created using the two BSTX1 restriction sites in the pJFE14 vector.

COS-7 cells were transiently transfected with either the pJFE14 library or control vector by the DEAE-dextran transfection protocol. Briefly, COS-7 cells were plated at a density of 1.0×10^6 cells/100 mm plate 24 hours prior to transfection. For transfection, the cells were cultured in serum-free DMEM containing 400 µg/ml of DEAE-dextran, 1 µM chloroquine, and 2 mM glutamine, and 1 µg of the appropriate DNA for 3-4 hours at 37°C in an atmosphere of 5% CO₂. The transfection media was aspirated and replaced with PBS with 10% DMSO for 2-3 min. Following this DMSO "shock", the COS-7 cells were placed into DMEM with 10% FBS, 1% each of penicillin and streptomycin, and 2 mM glutamine for 48 hours.

Because the TIE-2 ligand is secreted it was necessary to permeabilize the cells to detect binding of the receptorbody probe to the ligand. Two days after transfection the cells were rinsed with PBS and then incubated with PBS containing 1.8% formaldehyde for 15-30 min. at room temperature. Cells were then washed with PBS and incubated for 15 min. with PBS containing 0.1% Triton X-100 and 10% Bovine Calf Serum to permeabilize the cells and block non-specific binding sites.

The screening was conducted by direct localization of staining using a TIE-2 receptorbody (RB), which consisted of the extracellular domain of TIE-2 fused to the IgG1 constant region. This receptorbody was prepared as set forth in Example 2. A 100 mm dish of transfected, fixed and permeabilized COS cells was probed by incubating them for 30 min with TIE-2 RB. The cells were then washed twice with PBS and

incubated for an additional 30 min with PBS/10% Bovine Calf Serum/anti-human IgG-alkaline phosphatase conjugate. After three PBS washes, cells were incubated in alkaline-phosphatase substrate for 30-60 min. The dish was then inspected microscopically for the presence of stained cells. For each stained cell, a small area of cells including the stained cell was scraped from the dish using a plastic pipette tip and plasmid DNA was then rescued and used to electroporate bacterial cells. Single bacterial colonies resulting from the electroporation were picked and plasmid DNA prepared from these colonies was used to transfect COS-7 cells which were probed for TIE-2 ligand expression as evidenced by binding to TIE-2 receptorbodies. This allowed identification of single clones coding for TIE-2 ligand. Confirmation of TIE-2 ligand expression was obtained by phosphorylation of the TIE-2 receptor using the method set forth in Example 5. A plasmid clone encoding the TIE-2 ligand was deposited with the ATCC on October 7, 1994 and designated as "pJFE14 encoding TIE-2 ligand" under ATCC Accession No. 75910.

EXAMPLE 7 - ISOLATION AND SEQUENCING OF FULL LENGTH cDNA CLONE ENCODING HUMAN TIE-2 LIGAND

A human fetal lung cDNA library in lambda gt-10 (see Figure 3) was obtained from Clontech Laboratories, Inc. (Palo Alto, CA). Plaques were plated at a density of $1.25 \times 10^6/20 \times 20$ cm plate, and replica filters taken following standard procedures (Sambrook, et al., Molecular Cloning: A Laboratory Manual, 2nd Ed., page 8.46, Cold Spring Harbor Laboratory, Cold Spring Harbor, New York).

Isolation of human tie-2 ligand clones was carried out as follows. A 2.2 kb XhoI fragment from the deposited tie-2 ligand clone (ATCC NO.

75910 - see Example 6 above) was labeled by random priming to a specific activity of approximately 5×10^8 cpm/ng. Hybridization was carried out at 65°C in hybridization solution containing 0.5 mg/ml salmon sperm DNA. The filters were washed at 65°C in 2 x SSC, 0.1 % SDS and exposed to Kodak XAR-5 film overnight at -70°C. Positive phage were plaque purified. High titre phage lysates of pure phage were used for isolation of DNA via a Qiagen column using standard techniques (Qiagen, Inc., Chatsworth, CA, 1995 catalog, page 36). Phage DNA was digested with EcoRI to release the cloned cDNA fragment for subsequent subcloning. A lambda phage vector containing human tie-2 ligand DNA was deposited with the ATCC on October 26, 1994 under the designation λ gt10 encoding htie-2 ligand 1 (ATCC Accession No. 75928). Phage DNA may be subjected directly to DNA sequence analysis by the dideoxy chain termination method (Sanger, et al., 1977, Proc. Natl. Acad. Sci. U.S.A. 74: 5463-5467).

Subcloning of the human tie-2 ligand DNA into a mammalian expression vector may be accomplished as follows. The clone λ gt10 encoding htie-2 ligand 1 contains an EcoRI site located 490 base pairs downstream from the start of the coding sequence for the human TIE-2 ligand. The coding region may be excised using unique restriction sites upstream and downstream of the initiator and stop codons respectively. For example, an SpeI site, located 70 bp 5' to the initiator codon, and a Bpu1102i (also known as BlnI) site, located 265 bp 3' to the stop codon, may be used to excise the complete coding region. This may then be subcloned into the pJFE14 cloning vector, using the XbaI (compatible to the SpeI overhang) and the PstI sites (the PstI and Bpu1102i sites are both made blunt ended).

The coding region from the clone λ gt10 encoding htie-2 ligand 1 was sequenced using the ABI 373A DNA sequencer and Taq Dideoxy Terminator Cycle Sequencing Kit (Applied Biosystems, Inc., Foster City, CA). The nucleotide and deduced amino acid sequence of human TIE-2 ligand from the clone λ gt10 encoding htie-2 ligand 1 is shown in Figure 4.

In addition, full length human tie-2 ligand cDNA clones were obtained by screening a human glioblastoma T98G cDNA library in the pJFE14 vector. Clones encoding human TIE-2 ligand were identified by DNA hybridization using a 2.2 kb XhoI fragment from the deposited tie-2 ligand clone (ATCC NO. 75910) as a probe (see Example 6 above). The coding region was sequenced using the ABI 373A DNA sequencer and Taq Dideoxy Terminator Cycle Sequencing Kit (Applied Biosystems, Inc., Foster City, CA). This sequence was nearly identical to that of clone λ gt10 encoding htie-2 ligand 1. As shown in Figure 4, the clone λ gt10 encoding htie-2 ligand 1 contains an additional glycine residue which is encoded by nucleotides 1114-1116. The coding sequence of the T98G clone does not contain this glycine residue but otherwise is identical to the coding sequence of the clone λ gt10 encoding htie-2 ligand 1. Figure 5 sets forth the nucleotide and deduced amino acid sequence of human TIE-2 ligand from the T98G clone.

EXAMPLE 8 - ISOLATION AND SEQUENCING OF SECOND FULL LENGTH cDNA CLONE A ENCODING HUMAN TIE-2 LIGAND

A human fetal lung cDNA library in lambda gt-10 (see Figure 3) was obtained from Clontech Laboratories, Inc. (Palo Alto, CA). Plaques were plated at a density of 1.25×10^6 /20x20 cm plate, and replica

filters taken following standard procedures (Sambrook, et al.,
Molecular Cloning: A Laboratory Manual, 2nd Ed., page 8.46, Cold Spring
Harbor Laboratory, Cold Spring Harbor, New York). Duplicate filters
were screened at low stringency (2 x SSC, 55° C) with probes made to
5 the human TIE-2 ligand 1 sequence. One of the duplicate filters was
probed with a 5' probe, encoding amino acids 25 - 265 of human TIE-2
ligand 1 as set forth in Figure 4. The second duplicate filter was
probed with a 3' probe, encoding amino acids 282 - 498 of human TIE-2
ligand 1 sequence (see Figure 4). Both probes were hybridized at 55° C
10 in hybridization solution containing 0.5 mg/ml salmon sperm DNA.
Filters were washed in 2 x SSC at 55° C and exposed overnight to X-ray
film. In addition, duplicate filters were also hybridized at normal
stringency (2 x SSC, 65° C) to the full length coding probe of mouse
TIE-2 ligand 1 (F3-15, XhoI insert). Three positive clones were picked
15 that fulfilled the following criteria: i. hybridization had not been seen
to the full length (mouse) probe at normal stringency, and ii.
hybridization was seen at low stringency to both 5' and 3' probes.
EcoRI digestion of phage DNA obtained from these clones indicated two
independent clones with insert sizes of approximately 2.2kb and
20 approximately 1.8 kb. The 2.2kb EcoRI insert was subcloned into the
EcoRI sites of both pBluescript KS (Stratagene) and a mammalian
expression vector suitable for use in COS cells. Two orientations were
identified for the mammalian expression vector. The 2.2kb insert in
pBluescript KS was deposited with the ATCC on December 9, 1994 and
25 designated as pBluescript KS encoding human TIE 2 ligand 2. The start
site of the TIE-2 ligand 2 coding sequence is approximately 355 base
pairs downstream of the pBluescript EcoRI site.

COS-7 cells were transiently transfected with either the
expression vector or control vector by the DEAE-dextran transfection

protocol. Briefly, COS-7 cells were plated at a density of 1.0×10^6 cells/100 mm plate 24 hours prior to transfection. For transfection, the cells were cultured in serum-free DMEM containing 400 $\mu\text{g/ml}$ of DEAE-dextran, 1 μM chloroquine, and 2 mM glutamine, and 1 μg of the appropriate DNA for 3-4 hours at 37°C in an atmosphere of 5% CO_2 . The transfection media was aspirated and replaced with phosphate-buffered saline with 10% DMSO for 2-3 min. Following this DMSO "shock", the COS-7 cells were placed into DMEM with 10% FBS, 1% each of penicillin and streptomycin, and 2 mM glutamine for 48 hours.

Because the TIE-2 ligand is secreted it was necessary to permeabilize the cells to detect binding of the receptorbody probe to the ligand. Transfected COS-7 cells were plated at a density of 1.0×10^6 cells/100 mm plate. The cells were rinsed with PBS and then incubated with PBS containing 1.8% formaldehyde for 15-30 min. at room temperature. Cells were then washed with PBS and incubated for 15 min. with PBS containing 0.1% Triton X-100 and 10% Bovine Calf Serum to permeabilize the cells and block non-specific binding sites. The screening was conducted by direct localization of staining using a TIE-2 receptorbody, which consisted of the extracellular domain of TIE-2 fused to the IgG1 constant region. This receptorbody was prepared as set forth in Example 2. Transfected COS cells were probed by incubating them for 30 min with TIE-2 receptorbody. The cells were then washed twice with PBS, fixed with methanol, and then incubated for an additional 30 min with PBS/10% Bovine Calf Serum/anti-human IgG-alkaline phosphatase conjugate. After three PBS washes, cells were incubated in alkaline-phosphatase substrate for 30-60 min. The dish was then inspected microscopically for the presence of stained

cells. Cells expressing one orientation of the clone, but not the other orientation, were seen to bind the TIE-2 receptorbody.

One of skill in the art will readily see that the described methods may be used to further identify other related members of the TIE ligand family.

The coding region from the clone pBluescript KS encoding human TIE-2 ligand 2 was sequenced using the ABI 373A DNA sequencer and Taq Dideoxy Terminator Cycle Sequencing Kit (Applied Biosystems, Inc., Foster City, CA). The nucleotide and deduced amino acid sequence of human TIE-2 ligand 2 is shown in Figure 6.

EXAMPLE 9 - TIE-2 LIGAND 2 IS A RECEPTOR ANTAGONIST

Conditioned media from COS cells expressing either TIE-2 ligand 2 (TL2) or TIE-2 ligand 1 (TL1) were compared for their ability to activate TIE-2 receptors naturally present in human endothelial cell lines.

Lipofectamine reagent (GIBCO-BRL, Inc.) and recommended protocols were used to transfect COS-7 cells with either the pJFE14 expression vector alone, pJFE14 vector containing the human TIE-2 ligand 1 cDNA, or with a pMT21 expression vector (Kaufman, R.J., 1985, Proc. Natl. Acad. Sci. USA 82: 689-693) containing the human TIE-2 ligand 2 cDNA. COS media containing secreted ligands were harvested after three days and concentrated 20-fold by diafiltration (DIAFLO ultrafiltration membranes, Amicon, Inc.). The quantity of active TIE-2 ligand 1 and TIE-2 ligand 2 present in these media was determined and expressed as the amount (in resonance units, R.U.) of TIE-2 receptor specific binding activity measured by a BIAcore binding assay.

Northern (RNA) analyses revealed significant levels of TIE-2 transcripts in HAEC (Human Aortic Endothelial Cell) human primary endothelial cells (Clonetics, Inc.). Therefore, these cells were used to examine whether TIE-2 receptor is tyrosine-phosphorylated when
5 exposed to COS media containing the TIE-2 ligands. HAEC cells were maintained in a complete endothelial cell growth medium (Clonetics, Inc.) that contained 5% fetal bovine serum, soluble bovine brain extract, 10 ng/ml human EGF, 1 mg/ml hydrocortisone, 50 mg/ml gentamicin and 50 ng/ml amphotericin-B. Assessment of whether TL1 and TL2
10 could activate TIE-2 receptor in the HAEC cells was done as follows. Semi-confluent HAEC cells were serum-starved for two hours in high-glucose Dulbecco's MEM with added L-glutamine and penicillin-streptomycin at 37°C followed by replacement of the starvation medium with ligand-containing conditioned COS media for 7 minutes at
15 37°C in a 5% CO₂ incubator. The cells were subsequently lysed and TIE-2 receptor protein was recovered by immunoprecipitation of the lysates with TIE-2 peptide antiserum, followed by Western blotting with antiphosphotyrosine antiserum, exactly as described in example 1. The results are shown in Figure 7. Phosphotyrosine levels on the TIE-2
20 receptor (TIE-2-R) were induced by treatment of HEAC cells with TIE-2 ligand 1 (Lane L1) but not by TIE-2 ligand 2 (Lane L2) conditioned COS media. MOCK is conditioned media from COS transfected with JFE14 empty vector.

Evidence that both TL1 and TL2 specifically bind to the TIE-2
25 receptor was demonstrated by using a BIAcore to assay the TIE-2 receptor specific binding activities in transfected COS media and by immunostaining of TL1- and TL2-expressing COS cells with TIE-2 receptorbodies.

Because TL2 did not activate the TIE-2 receptor, applicants set out to determine whether TL2 might be capable of serving as an antagonist of TL1 activity. HAEC phosphorylation assays were performed in which cells were first incubated with an "excess" of TL2, followed by addition of dilute TL1. It was reasoned that prior occupancy of TIE-2 receptor due to high levels of TL2 might prevent subsequent stimulation of the receptor following exposure to TL1 present at a limiting concentration.

Semi-confluent HAEC cells were serum-starved as described above and then incubated for 3 min., at 37°C with 1-2 ml. of 20X COS/JFE14-TL2 conditioned medium. Control plates were treated with 20X COS/JFE14-only medium (MOCK). The plates were removed from the incubator and various dilutions of COS/JFE14-TL1 medium were then added, followed by further incubation of the plates for 5-7 min. at 37°C. Cells were subsequently rinsed, lysed and TIE-2-specific tyrosine phosphorylation in the lysates was examined by receptor immunoprecipitation and Western blotting, as described above. TL1 dilutions were made using 20X COS/JFE14-TL1 medium diluted to 2X, 0.5X, 0.1X, or 0.02X by addition of 20X COS/JFE14-alone medium. An assay of the initial 20X TL1 and 20X TL2 COS media using BIAcore biosensor technology indicated that they contained similar amounts of TIE-2-specific binding activities, i.e., 445 R.U. and 511 R.U. for TL1 and TL2, respectively. The results of the antiphosphotyrosine Western blot, shown in Figure 8, indicate that when compared to prior treatment of HAEC cells with MOCK medium (lane 1), prior treatment of HAEC cells with excess TIE-2 ligand 2 (lane 2) antagonizes the subsequent ability of dilute TIE-2 ligand 1 to activate the TIE-2 receptor (TIE-2-R).

The ability of TL2 to competitively inhibit TL1 activation of the TIE-2-R was further demonstrated using the human cell hybrid line,

EA.hy926 (see Example 21 for detailed description of this cell line and its maintenance). Experiments were performed in which unconcentrated COS cell media containing TL1 were mixed at varying dilutions with either MOCK- or TL2- conditioned media and placed on serum-starved EA.hy926 cell monolayers for 5 minutes at 37°C. The media were then removed, the cells were harvested by lysis and TIE-2-specific tyrosine phosphorylation was examined by Western blots, as described above. Figure 9 shows an experiment which contains three groups of treatments, as viewed from left to right. As shown in the four lanes at the left, treatment of the EA.hy926 cells with 1x COS-TL1 alone robustly activated the endogenous TIE-2-R in these cells, whereas 1x TL2 COS medium was inactive. However, mixture of TL1 with either MOCK or TL2 demonstrated that TL2 can block the activity of TL1 in a dose-dependent fashion. In the central three pairs of lanes the ratio of TL2 (or MOCK) was decreased while the amount of TL1 in the mixture was correspondingly increased from 0.1x to 0.3x. At any of these mixture ratios the TL1:TL2 lanes showed a reduced level of TIE-2-R phosphorylation compared to that of the corresponding TL1:MOCK lanes. When the amount TL1 was held steady and the amount of TL2 (or MOCK) was decreased, however (shown in the three pairs of lanes at the right), a point was reached at which the TL2 in the sample was too dilute to effectively inhibit TL1 activity. The relative amount of each ligand present in these conditioned COS media could be estimated from their binding units as measured by the BIAcore assay and from Western blots of the COS media with ligand-specific antibodies. Consequently, we can infer that only a few-fold molar excess of TL2 is required to effectively block the activity of TL1 in vitro. This is significant because we have observed distinct examples in vivo (see Example 17 and Figure 16) where TL2 mRNAs achieve considerable abundance

relative to those of TL1. Thus, TL2 may be serving an important physiological role in effectively blocking signaling by the TIE-2-R at these sites.

Taken together these data confirm that, unlike TL1, TL2 is unable
5 to stimulate endogenously expressed TIE-2-R on endothelial cells. Furthermore, at a few fold molar excess TL2 can block TL1 stimulation of the TIE-2 receptor, indicating that TL2 is a naturally occurring TIE-2 receptor antagonist.

10 **EXAMPLE 10 - IDENTIFICATION OF TIE-2-SPECIFIC BINDING ACTIVITY**
IN CONDITIONED MEDIUM AND COS CELL SUPERNATANTS

Binding activity of 10x CCM from the cell lines C2C12-ras, Rat2
ras, SHEP, and T98G, or COS cell supernatants after transfection with
15 either human TIE-2 ligand 1 (hTL1) or human TIE-2 ligand 2 (hTL2) was measured using biosensor technology (BIAcore; Pharmacia Biosensor, Piscataway, NJ) which monitors biomolecular interactions in real-time via surface plasmon resonance (SPR). Purified rat or human TIE-2 RB
was covalently coupled through primary amines to the carboxymethyl
20 dextran layer of a CM5 research grade sensor chip (Pharmacia Biosensor; Piscataway, NJ). The sensor chip surface was activated using a mixture of N-hydroxysuccinimide (NHS) and N-ethyl-N'-(3-dimethylaminopropyl)carbodiimide (EDC), followed by immobilization of TIE-2 RB (25 µg/mL, pH 4.5) and deactivation of unreacted sites with
25 1.0 M ethanolamine (pH 8.5). In general, 9000-10000 RU of each receptorbody was coupled to the sensor chip.

The running buffer used in the system was HBS (10 mM Hepes, 150 mM NaCl, 0.005% P20 surfactant, pH 7.4). The samples were

centrifuged for 15 min at 4°C and further clarified using a sterile, low protein-binding 0.45 µm filter (Millipore; Bedford, MA). Dextran (2mg/ml) and P20 surfactant (0.005%) were added to each sample. Aliquots of 40 µL were injected across the immobilized surface (either
5 rat or human TIE-2) at a flow rate of 5 µL/min and the receptor binding was monitored for 8 min. The binding activity (resonance units, RU) was measured as the difference between a baseline value determined 30 s prior to the sample injection and a measurement taken at 30 s post-injection. Regeneration of the surface was accomplished with
10 one 15-µL pulse of 3 M MgCl₂.

The CCM samples (C2C12-ras, Rat2-ras, SHEP, T98G) were tested on the rat TIE-2 RB immobilized surface, while the recombinant hTL1 and hTL2 were tested on the human TIE-2 RB immobilized surface. In each case, specific binding to the TIE-2 receptorbody was evaluated by
15 incubating the samples with 25 µg/ml of either soluble TIE-2 (rat or human) RB or trkB RB prior to assaying the binding activity. As shown in Figures 10 and 11, the addition of soluble trkB RB causes a slight decrease in the TIE-2 binding activity, while the addition of soluble TIE-2 RB significantly reduces the binding activity as compared to that
20 measured in the absence of TIE-2 RB.

EXAMPLE 11 - TIE-2 RB SPECIFICALLY BLOCKS ACTIVATION OF THE TIE-2 RECEPTOR BY TIE-2 LIGAND 1

25 The applicants sought to determine whether soluble TIE-2 RB can serve as a competitive inhibitor to block activation of TIE-2 receptor by TIE-2 ligand 1 (TL1). To do this, TL1-containing COS media were preincubated with either TIE-2- or TrkB-RB and then compared for

their ability to activate TIE-2 receptors naturally present in a human endothelial cell line.

Conditioned COS media were generated from COS-7 cells transfected with either the pJFE14 expression vector alone (MOCK), or
5 pJFE14 vector containing the human TIE-2 ligand 1 cDNA (TL1) and harvested as described in Example 9 hereinabove, with the exception that the media were sterile filtered but not concentrated. The quantity of TL1 was determined and expressed as the amount (in resonance units, R.U.) of TIE-2 receptor-specific binding activity measured by
10 BIAcore binding assay.

Northern (RNA) analyses revealed significant levels of tie-2 transcripts in HUVEC (Human Umbilical Vein Endothelial Cell) human primary endothelial cells (Clonetics, Inc.). Therefore, these cells were used to examine whether TIE-2 receptor can be tyrosine-
15 phosphorylated when exposed in the presence of TIE-2- or TrkB-RBs to COS media containing TL1. HUVEC cells were maintained at 37°C, 5% CO₂ in a complete endothelial cell growth medium (Clonetics, Inc.) that contained 5% fetal bovine serum, soluble bovine brain extract with 10 µg/ml heparin, 10 ng/ml human EGF, 1 µg/ml hydrocortisone, 50 µg/ml
20 gentamicin and 50 ng/ml amphotericin-B. Assessment of whether TL1 could activate TIE-2 receptor in the HUVEC cells was done as follows. Confluent dishes of HUVEC cells were serum-starved for two-to-four hours in low-glucose Dulbecco's MEM at 37°C, 5% CO₂, followed by 10 minute incubation in starvation medium that included 0.1 mM sodium
25 orthovanadate, a potent inhibitor of phosphotyrosine phosphatases. Meanwhile, conditioned COS media were preincubated 30 min. at room temperature with either TIE-2- or TrkB-RB added to 50 µg/ml. The starvation medium was then removed from the HUVEC dishes and

incubated with the RB-containing COS media for 7 minutes at 37°C. HUVEC cells were subsequently lysed and TIE-2 receptor protein was recovered by immunoprecipitation with TIE-2 peptide antiserum, followed by Western blotting with an anti-phosphotyrosine antibody, as described in Example 1. The results are shown in Figure 12.

Phosphotyrosine levels on the TIE-2 receptor were induced by treatment of HUVEC cells with TIE-2 ligand 1 (TL1) relative to that seen with control medium (MOCK) and this induction is specifically blocked by prior incubation with TIE-2-RB (TIE-2-Fc) but not by incubation with TrkB-RB (TrkB-Fc). These data indicate that soluble TIE-2 RB can serve as a selective inhibitor to block activation of TIE-2 receptor by TIE-2 ligand 1.

EXAMPLE 12 - CONSTRUCTION OF TIE-2 LIGANDBODIES

An expression construct was created that would yield a secreted protein consisting of the entire coding sequence of human TIE-2 ligand 1 (TL1) or TIE-2 ligand 2 (TL2) fused to the human immunoglobulin gamma-1 constant region (IgG1 Fc). These fusion proteins are called TIE-2 "ligandbodies" (TL1-Fc or TL2-Fc). The Fc portion of TL1-Fc and TL2-Fc was prepared as follows. A DNA fragment encoding the Fc portion of human IgG1 that spans from the hinge region to the carboxy-terminus of the protein, was amplified from human placental cDNA by PCR with oligonucleotides corresponding to the published sequence of human IgG1; the resulting DNA fragment was cloned in a plasmid vector. Appropriate DNA restriction fragments from a plasmid encoding full-length TL1 or TL2 and from the human IgG1 Fc plasmid were ligated on either side of a short PCR-derived fragment that was designed so as to

fuse, in-frame, TL1 or TL2 with human IgG1 Fc protein-coding sequences.

Milligram quantities of TL2-Fc were obtained by cloning the TL2-Fc DNA fragment into the pVL1393 baculovirus vector and subsequently infecting the *Spodoptera frugiperda* SF-21AE insect cell line.

5 Alternatively, the cell line SF-9 (ATCC Accession No. CRL-1711) or the cell line BTI-TN-5b1-4 may be used. DNA encoding the TL2-Fc was cloned as an Eco RI-NotI fragment into the baculovirus transfer plasmid pVL1393. Plasmid DNA was recombined into viral DNA by mixing 3 µg of plasmid DNA with 0.5 µg of Baculo-Gold DNA (Pharminigen), followed
10 by introduction into liposomes using 30µg Lipofectin (GIBCO-BRL).

DNA-liposome mixtures were added to SF-21AE cells (2x 10⁶ cells/60mm dish) in TMN-FH medium (Modified Grace's Insect Cell Medium (GIBCO-BRL) for 5 hours at 27°C, followed by incubation at 27°C for 5 days in TMN-FH medium supplemented with 5% fetal calf
15 serum. Tissue culture medium was harvested for plaque purification of recombinant viruses, which was carried out using methods previously described (O'Reilly, D.R., L.K. Miller, and V.A. Luckow, Baculovirus Expression Vectors - A Laboratory Manual. 1992, New York: W.H.

Freeman) except that the agarose overlay contained 125 mg/mL X-gal
20 (5-bromo-4-chloro-3-indolyl-b- D-galactopyranoside; GIBCO-BRL).

After 5 days of incubation at 27°C, non-recombinant plaques were scored by positive chromogenic reaction to the X-gal substrate, and their positions marked. Recombinant plaques were then visualized by addition of a second overlay containing 100 mg/mL MTT (3-[4,5-
25 dimethylthiazol-2-yl]2,5-diphenyltetrazolium bromide; Sigma).

Putative recombinant virus plaques were picked by plug aspiration, and purified by multiple rounds of plaque isolation to assure homogeneity. Virus stocks were generated by serial, low-multiplicity passage of

plaque-purified virus. Low passage stocks of one virus clone (vTL2-Fc Clone #7) were produced.

SF-21AE cells were cultured in serum-free medium (SF-900 II, Gibco BRL) containing 1X antibiotic/antimycotic solution (Gibco BRL) and 25 mg/L Gentamycin (Gibco BRL). Pluronic F-68 was added as a surfactant to a final concentration of 1g/L. Cultures (4L) were raised in a bioreactor (Artisan Cell Station System) for at least three days prior to infection. Cells were grown at 27°C, with gassing to 50 % dissolved oxygen, at a gas flow rate of 80 mL/min (aeration at a sparge ring). Agitation was by means of a marine impeller at a rate of 100 rpm. Cells were harvested in mid-logarithmic growth phase ($\sim 2 \times 10^6$ cells/mL), concentrated by centrifugation, and infected with 5 plaque forming units of vTL2-Fc per cell. Cells and inoculum were brought to 400mL with fresh medium, and virus was adsorbed for 2 hours at 27°C in a spinner flask. The culture was then resuspended in a final volume of 8L with fresh serum-free medium, and the cells incubated in the bioreactor using the previously described conditions.

Culture medium from vTL2-Fc-infected SF21AE cells were collected by centrifugation (500x g, 10 minutes) at 72 hours post-infection. Cell supernatants were brought to pH 8 with NaOH. EDTA was added to a final concentration of 10 mM and the supernatant pH was readjusted to 8. Supernatants were filtered (0.45 μ m, Millipore) and loaded on a protein A column (protein A sepharose 4 fast flow or HiTrap protein A, both from Pharmacia). The column was washed with PBS containing 0.5 M NaCl until the absorbance at 280 nm decreased to baseline. The column was washed in PBS and eluted with 0.5 M acetic acid. Column fractions were immediately neutralized by eluting into

tubes containing 1 M Tris pH 9. The peak fractions containing the TL2-Fc were pooled and dialyzed versus PBS.

EXAMPLE 13- EXPRESSION OF TIE-1, TIE-2, TL1, AND TL2 IN RENAL CELL CARCINOMA

In situ hybridization experiments were performed on human renal cell carcinoma tumor tissue using TIE-1, TIE-2, TL1, and TL2 cDNA probes. TIE-2, TIE-1, TL1, and TL2 expression were all up-regulated in the tumor vasculature. Ligand expression appeared to be localized to either the vascular endothelial cells (TL2) or very near the vascular endothelial cells in the mesenchyme (TL1). VEGF has been shown to be dramatically up-regulated in this tumor tissue. Brown, et al. Am. J. Pathol. 143:1255-1262 (1993).

EXAMPLE 14- EXPRESSION OF TIE-1, TIE-2, TL1, AND TL2 IN WOUND HEALING

In situ hybridization experiments were performed on cross-sectional tissue slices obtained from a rat cutaneous wound model using TIE-1, TIE-2, TL1, and TL2 cDNA probes. The wound healing model involves pressing a small cork bore against the skin of a rat and removing a small, cylindrical plug of skin. As healing begins at the base of the wound, a vertical slice of tissue is taken and used for *in situ* hybridization. In the tested tissue sample, TL1 and TL2 appeared to be slightly up-regulated by four days post-injury. In contrast to the slightly up-regulated expression of TL1 and TL2 in this tissue, VEGF expression, which may precede TL1 and TL2 expression, is dramatically up-regulated.

EXAMPLE 15- EXPRESSION OF TIE LIGANDS IN FETAL LIVER AND THYMUS

Reverse transcription-PCR (RT-PCR) was performed on mouse
5 E14.5 fetal liver and mouse E17.5 fetal thymus. Agarose gel
electrophoresis of the RT-PCR products revealed that in the mouse
fetal liver, TIE-2 ligand 1 (TL1) RNA is enriched in the stromal region,
but is absent in c-kit⁺TER119 hematopoietic precursor cells. In this
same tissue, TIE-2 ligand 2 (TL2) RNA is enriched in the stromal cells,
10 but absent in the hematopoietic precursor cells (Figure 13). In the
mouse fetal thymus, TL2 is enriched in the stromal cells (Figure 14).

EXAMPLE 16- THE TIE RECEPTOR/LIGAND SYSTEM IN ANGIOGENESIS

15

Although the TIE-2/TIE ligand system appears to play an
important role in endothelial cell biology, it has not been shown to play
a significant, active role in the early to intermediate stages of
vascularization (e.g. angioblast or endothelial cell proliferation and
20 migration, tubule formation, and other early stage events in vascular
modeling). In contrast to the receptors and factors known to mediate
these aspects of vascular development, the temporally late pattern of
expression of TIE-2 and TL1 in the course of vascularization suggests
that this system plays a distinct role in the latter stages vascular
25 development, including the structural and functional differentiation
and stabilization of new blood vessels. The pattern of expression of
TIE-2/TL1 also is consistent with a continuing role in the maintenance
of the structural integrity and/or physiological characteristics of an
established vasculature.

TIE Ligand 2 (TL2) appears to be a competitive inhibitor of TL1. The spatiotemporal characteristics of TL2 expression suggest that this single inhibitory molecule may play multiple, context-dependent roles essential to appropriate vascular development or remodeling (e.g. de-
5 stabilization/de-differentiation of mature endothelial cells allowing the formation of new vessels from existing vasculature, inhibition of inappropriate blood vessel formation, and regression/involution of mature blood vessels). Figure 15 is a schematic representation of the hypothesized role of the TIE-2/TIE ligands in angiogenesis. In this
10 figure TL1 is represented by (•), TL2 is represented by (*), TIE-2 is represented by (T), VEGF is represented by ([]), and flk-1 (a VEGF receptor) is represented by (Y).

15 EXAMPLE 17- EXPRESSION OF TIE LIGANDS IN THE FEMALE
REPRODUCTIVE SYSTEM: EXPRESSION IN THE
OVARY

Preliminary observations made in experiments examining the expression of the TIE receptors and ligands in the female reproductive
20 system are consistent with the hypothesis the TL1 plays a role in neovascularization which temporally follows that of VEGF. The pattern of TL2 expression is also consistent with an antagonism of the action of TL1, and a specific role in vascular regression. To verify this, expression of relevant mRNAs can be examined following experimental
25 induction of follicular and luteal development so that their temporal relation to various aspects of neovascularization/vascular regression can be more clearly defined (e.g. in conjunction with endothelial cell staining, vascular fills). Angiogenesis associated with follicular development and corpus luteum formation in staged ovaries of mature,
30 female rats or following induced ovulation in pre-pubertal animals was

followed using *in situ* hybridization. Figure 16 contains photographs of *in situ* hybridization slides showing the temporal expression pattern of TIE-2, TL1, TL2, and VEGF during the ovarian cycle [Column 1: Early pre-ovulatory follicle; Column 2: pre-ovulatory follicle; Column 3: early corpus luteum; and Column 4: atretic follicle; Row A: bright field; Row B: VEGF; Row C: TL2; Row D: TL1 and Row E: TIE-2 receptor]. These studies revealed that VEGF, TL1 and TL2 are expressed in a temporally and spatially coordinate fashion with respect to the development and regression of vasculature in the ovary, specifically with respect to the establishment of the vascular system which is generated in the course of the conversion of an ovarian follicle to a corpus luteum (CL).

Briefly, VEGF expression increases in the follicular granule layer prior to its vascularization during the process of luteinization. During the process of CL formation, highest levels of VEGF expression are apparent in the center of the developing CL in the vicinity of luteinizing cells which are not yet vascularized. VEGF levels remain moderately high and are diffusely distributed in the developed CL. In contrast, noticeably enhanced expression of TIE-2 ligand 1 occurs only late in process of CL formation, after a primary vascular plexus has been established. Later, TL1 expression is apparent throughout the CL at which time the definitive capillary network of the CL has been established.

TL2 exhibits a more complex pattern of expression than either VEGF or TL1. In the developing CL, TL2 is expressed at highest levels at the front of the developing capillary plexus- between the central avascular region of the CL where VEGF expression is highest, and the most peripheral portion of the CL where TL1 expression is dominant and where the luteinization process is complete and the vascular system is most mature. TL2 also appears to be expressed at high levels in the

follicular layer of large follicles which are undergoing atresia. While TL1 is also apparent in atretic follicles, VEGF is not expressed.

The pattern of expression described above is most consistent with a role for VEGF in the initiation of angiogenesis, with TL1 acting late in this process-for example in modeling and/or stabilization of the definitive vascular network. In contrast, TL2 is present both in areas of active expansion of a newly forming vascular network (during CL formation), and in regions which fail to establish a new vasculature and vascular regression is in progress (atretic follicles). This suggests a more dynamic and complex role for TL2, possibly involving destabilization of existing vasculature (necessary for regression) or developing vasculature (necessary for the dynamic modeling of newly forming vessels).

15 EXAMPLE 18- CONSTRUCTION AND CHARACTERIZATION OF
THE CYS-TL1 MUTANT

The TIE-2 ligands have two major structural domains, one described as a "coiled-coil" domain comprising the approximate C-terminal third of the protein and the other a "fibrinogen-like" domain comprising the approximate N-terminal two-thirds of the protein. Although the TIE-2 ligands, designated TL1 and TL2, share similar structural homology, they exhibit different physical and biological properties. Under non-reducing electrophoretic conditions, both proteins exhibit covalent, multimeric structures, with TL1 existing primarily as a trimer and TL2 existing primarily as a dimer. Figure 17 is a schematic representation of how the TIE-2 ligands may be interacting to form multimers. In terms of biological activity, TL1 has been shown to be an agonist of the TIE-2 receptor, as demonstrated by

induction of phosphorylation in TIE-2 expressing cells. TL2, on the other hand, appears to be a competitive inhibitor of TL1. Investigations into what factors might be contributing to the different physical and biological properties of the two molecules revealed the presence of a cysteine residue (CYS265) preceding the fibrinogen-like domain in TL1 but absent in TL2. This CYS265 residue in TL1 is encoded by TGC and is located at about nucleotides 1102-1104 at the approximate junction between the coiled-coil and fibrinogen-like domains. Because cysteine residues are generally involved in disulfide bond formation, the presence of which can contribute to both the tertiary structure and biological properties of a molecule, it was thought the perhaps the presence of the CYS265 in TL1 might be at least partially responsible for the different properties of the two molecules. To test this hypothesis, an expression plasmid was constructed which contained a mutation in TL1 in which the CYS was replaced with an amino acid which does not form disulfide bonds. In addition to this TL1/CYS⁻ mutant, a second expression plasmid was constructed which mutated the corresponding position in TL2 so that this residue was now a cysteine. Both non-mutated and mutated expression plasmids of TL1 and TL2 were transiently transfected into COS cells. Cell supernatants containing the recombinant proteins were harvested and samples subjected to both reducing and non-reducing SDS/PAGE electrophoresis and subsequent western blotting. Figure 18 contains western blots of both non-mutated and mutated TL1 and TL2 proteins under revealing that the TL1/CYS⁻ mutant behaves more TL2-like in that it runs as a dimer and that the TL2/CYS⁺ mutant behaves more TL1-like in that it is able to form a trimer as well as higher-order multimers. Interestingly, when the two mutant proteins were tested for their ability to induce phosphorylation in TIE-2 expressing cells, the TL1/CYS⁻ mutant was

able to activate the TIE-2 receptor, whereas the TL2/CYS⁺ mutant did not gain any activating activity.

**EXAMPLE 19- CONSTRUCTION AND CHARACTERIZATION OF
FIBRINOGEN-LIKE DOMAIN ONLY MUTANTS**

In order to test whether the fibrinogen-like domain (F-domain) of the TIE-2 ligands contained TIE-2 activating activity, expression plasmids were constructed which deleted the coiled-coil domain, leaving only that portion of the DNA sequence encoding the F-domain (beginning at about nucleotide 1159, amino acid residue ARG284). This mutant construct was transiently transfected into COS cells. The supernatant containing the recombinant protein was harvested. The TL1/F-domain mutant was tested for its ability to bind the TIE-2 receptor. The results showed that, as a monomer, the TL1/F-domain mutant was not able to bind TIE-2 at a detectable level. However, when the TL1/F-domain monomer was myc-tagged and subsequently clustered with an antibody directed against the myc tag, it did exhibit detectable binding to TIE-2. However, the antibody-clustered TL1/F-domain mutant was not able to induce phosphorylation in a TIE-2 expressing cell line. Figure 17 shows a schematic representation of the F-domain construct and its binding ability plus and minus antibody clustering.

**EXAMPLE 20- A RECEPTORBODY BINDING ASSAY AND A LIGAND
BINDING AND COMPETITION ASSAY**

A quantitative cell-free binding assay with two alternate formats has been developed for detecting either TIE-2 receptorbody binding or ligand binding and competition. In the receptorbody binding

version of the assay, TIE-2 ligands (purified or partially purified; either TL1 or TL2) are coated onto an ELISA plate. Receptorbody at varying concentrations is then added, which binds to the immobilized ligand in a dose-dependent manner. At the end of 2 hours, excess
5 receptorbody is washed away, then the amount bound to the plate is reported using a specific anti-human Fc antibody which is alkaline phosphatase tagged. Excess reporter antibody is washed away, then the AP reaction is developed using a colored substrate. The assay is quantitated using a spectrophotometer. Figure 19 shows a typical TIE-
10 2-IgG binding curve. This assay has been used to evaluate the integrity of TIE-2-IgG after injection into rats and mice. The assay can also be used in this format as a ligand competition assay, in which purified or partially-purified TIE ligands compete with immobilized ligand for receptorbody. In the ligand binding and competition version of the
15 binding assay, TIE-2 ectodomain is coated onto the ELISA plate. The Fc-tagged fibrinogen-like domain fragments of the TIE ligands (TL1-fFc and TL2-fFc) then bind to the ectodomain, and can be detected using the same anti-human Fc antibody as described above. Figure 20 shows an example of TL1-fFc binding to TIE-2 ectodomain. This version of the
20 assay can also be used to quantitate levels of TL1-fFc in serum or other samples. If untagged ligand (again, either purified or unpurified) is added at the same time as the TL1-fFc, then a competition is set up between tagged ligand fragment and full-length ligand. The full-length ligand can displace the Fc-tagged fragment, and a competition curve is
25 generated.

EXAMPLE 21- EA.hy926 CELL LINE CAN BE USED AS A REPORTER
CELL LINE FOR TIE LIGAND ACTIVITY

EA.hy926 is a cell hybrid line that was established by fusion of HUVEC with the human lung carcinoma-derived line, A549 [Edgell, et al. Proc. Natl. Acad. Sci. (USA) 80, 3734-3737 (1983). EA.hy926 cells have been found to express significant levels of TIE-2 receptor protein with low basal phosphotyrosine levels. The density at which EA.hy926 cells are passaged prior to their use for receptor assays, as well as their degree of confluency at the time of assay, can affect TIE-2 receptor abundance and relative inducibility in response to treatment with ligand. By adopting the following regimen for growing these cells the EA.hy926 cell line can be used as a dependable system for assay of TIE-2 ligand activities.

EA.hy926 cells are seeded at 1.5×10^6 cells in T-75 flasks (Falconware) and re-fed every other day with high-glucose Dulbecco's MEM, 10% fetal bovine serum, L-glutamine, penicillin-streptomycin, and 1x hypoxanthine-aminopterin-thymidine (HAT, Gibco/BRL). After three to four days of growth, the cells are passaged once again at 1.5×10^6 cells per T-75 flask and cultured an additional three to four days. For phosphorylation assays, cells prepared as described above were serum-starved by replacement of the culture medium with high-glucose DMEM and incubation for 2-3 hours at 37°C. This medium was aspirated from the flask and samples of conditioned media or purified ligand were added to the flask in a total volume of 1.5 ml followed by incubation at 37°C for 5 minutes. Flasks were removed from the incubator and placed on a bed of ice. The medium was removed and replaced with 1.25 ml Lysis Buffer containing 1% nonidet P-40, 0.5% sodium deoxycholate, 0.1% SDS in 20 mM Tris, pH 7.6, 150 mM NaCl, 50 mM NaF, 1mM sodium orthovanadate, 5 mM benzamidine, and 1mM EDTA containing the protease inhibitors PMSF, aprotinin, and leupeptin. After 10 minutes on

ice to allow membrane solubilization, plates were scraped and cell lysates were clarified by microcentrifugation at top speed for 10 minutes at 4°C. TIE-2 receptor was immunoprecipitated from the clarified supernatant by incubation in the cold with an anti-TIE-2 polyclonal antiserum and Protein G-conjugated Sepharose beads. The beads were washed three times with cold cell lysis buffer and boiled 5 minutes in Laemmli sample buffer, which was then loaded on 7.5% SDS-polyacrylamide gels. Resolved proteins were electrotransferred to PVDF (Lambdia-P) membrane and then subjected to Western blot analysis using anti-phosphotyrosine antibody and the ECL reagent. Subsequent comparison of total TIE-2 protein levels on the same blots was done by stripping the anti-phosphotyrosine antibody and reincubating with a polyclonal antiserum specific to the ectodomain of TIE-2.

DEPOSITS

The following have been deposited with the American Type Culture Collection, 12301 Parklawn Drive, Rockville, Maryland 20852 in accordance with the Budapest Treaty. A plasmid clone encoding a TIE-2 ligand was deposited with the ATCC on October 7, 1994 and designated as "pJFE14 encoding TIE-2 ligand" under ATCC Accession No. 75910. Recombinant Autographa californica baculovirus encoding TIE-2 receptorbody was deposited with the ATCC on October 7, 1994 and designated as "vTIE-2 receptorbody" under ATCC Accession No. VR2484. A lambda phage vector containing human tie-2 ligand DNA was deposited with the ATCC on October 26, 1994 and designated as λ gt10 encoding htie-2 ligand 1 under ATCC Accession No. 75928. A plasmid clone encoding a second TIE-2 ligand was deposited with the ATCC on

December 9, 1994 and designated as "pBluescript KS encoding human TIE 2 ligand 2" under ATCC Accession No. 75963.

The present invention is not to be limited in scope by the specific
5 embodiments described herein. Indeed, various modifications of the invention in addition to those described herein will become apparent to those skilled in the art from the foregoing description and accompanying figures. Such modifications are intended to fall within the scope of the appended claims.

CLAIMS

1. An isolated nucleic acid molecule encoding a TIE-2 ligand.
2. An isolated nucleic acid molecule according to claim 1 wherein the nucleic acid sequence is:
 - (a) the nucleic acid sequence comprising the coding region of the human TIE-2 ligand as set forth in Figure 4 or Figure 5;
 - (b) a nucleic acid sequence that hybridizes under moderately stringent conditions to the nucleic acid sequence of (a) and which encodes a TIE-2 ligand that binds TIE-2 receptor; or
 - (c) a nucleic acid sequence which, but for the degeneracy of the genetic code would hybridize to a nucleic acid sequence of (a) or (b), and which encodes a TIE-2 ligand that binds TIE-2 receptor.
3. An isolated nucleic acid molecule according to claim 2 wherein the encoded TIE-2 ligand is a TIE-2 agonist.
4. An isolated nucleic acid molecule according to claim 1 wherein the nucleic acid sequence is
 - (a) the nucleic acid sequence comprising the coding region of the human TIE-2 ligand as set forth in Figure 6;
 - (b) the nucleic acid sequence comprising the coding region of the fibrinogen-like domain of the human TIE-2 ligand as set forth in Figure 4, 5 or 6;
 - (c) a nucleic acid sequence that hybridizes under moderately stringent conditions to the nucleic acid sequence of (a) and which encodes a TIE-2 ligand that binds TIE-2 receptor; or

- (d) a nucleic acid sequence which, but for the degeneracy of the genetic code would hybridize to a nucleic acid sequence of (a) or (b), and which encodes a TIE-2 ligand that binds TIE-2 receptor
5. An isolated nucleic acid molecule according to claim 4 wherein the encoded TIE-2 ligand is a TIE-2 antagonist.
6. A vector which comprises a nucleic acid molecule of any one of the preceding claims.
7. A vector according to claim 6 wherein the nucleic acid molecule is operatively linked to an expression control sequence capable of directing its expression in a host cell.
8. A vector according to claim 6 or 7 which is a plasmid.
9. A plasmid according to claim 8 designated pJFE14 encoding TIE-2 ligand (ATCC Accession No. 75910).
10. A plasmid according to claim 8 designated pBluescript KS encoding human TIE-2 ligand 2 (ATCC Accession No. 75963).
11. A vector according to claim 6 or 7 designated as λ gt10 encoding hTIE2 ligand 1 (ATCC Accession No. 75928).
12. An isolated TIE-2 ligand substantially free of other proteins.
13. An isolated TIE-2 ligand according to claim 12 encoded by a nucleic acid molecule according to claim 1.

14. An isolated TIE-2 ligand according to claim 12 encoded by a nucleic acid according to claim 2 or 3.
15. An isolated TIE-2 ligand according to claim 12 encoded by a nucleic acid according to claim 4 or 5.
16. A host-vector system for the production of a ligand according to any one of claims 12 to 15 which comprises a vector according to any one of claims 6 to 11 in a host cell.
17. A host-vector system according to claim 16 wherein the host cell is a bacterial, yeast, insect or mammalian cell.
18. A host vector system comprising the host vector system of claim 16 or 17 and a nucleic acid encoding the TIE-2 receptor.
19. A method of producing a ligand as defined in any one of claims 12 to 15 which comprises growing cells of a host-vector system according to any one of claims 16 to 18 under conditions permitting production of the ligand, and recovering the ligand so produced.
20. An antibody which specifically binds the ligand of any one of claims 12 to 15.
21. An antibody according to claim 20 which is a monoclonal antibody.
22. A receptorbody which specifically binds the ligand of any one of claims 12 to 15.

23. An isolated nucleic acid molecule encoding a receptorbody according to claim 22.
24. A vector comprising a nucleic acid molecule according to claim 23.
25. A vector according to claim 24 which is a plasmid.
26. A plasmid according to claim 25 designated vTIE-2 receptorbody (ATCC Deposit VR2484).
27. A conjugate comprising a ligand according to any one of claims 12 to 15 and, conjugated thereto, a cytotoxic agent.
28. A conjugate according to claim 27 wherein the cytotoxic agent is a radioisotope or toxin.
29. A pharmaceutical composition comprising a TIE-2 ligand according to any one of claims 12 to 15 and a pharmaceutically acceptable carrier.
30. A pharmaceutical composition comprising an antibody according to claim 20 or 21 and a pharmaceutically acceptable carrier.
31. A pharmaceutical composition comprising a receptorbody according to claim 22 and a pharmaceutically acceptable carrier.
32. A pharmaceutical composition comprising a conjugate according to claim 27 or 29 and a pharmaceutically acceptable carrier.

33. A ligand according to any one of claims 12 to 15, an antibody according to claim 20 or 21, a receptorbody according to claim 22, a conjugate according to claim 27 or 29, or a composition according to any one of claims 29 to 32 for use in a method of treatment of the human or animal body, or in a method of diagnosis.
34. A ligand according to claim 14 for use in a method of treatment of the human or animal body.
35. A ligand according to claim 15 for use in a method of treatment of the human or animal body.
36. An antibody or receptorbody according to claim 33, in which antibody or receptorbody specifically binds the ligand of claim 14, for use in a method of blocking blood vessel growth in a mammal.
37. An antibody or receptorbody according to claim 36 for use in a method wherein the mammal is a human.
38. A ligand according to claim 34 for use in a method of promoting neovascularization in a mammal.
39. A ligand according to claim 38 for use in the promotion of wound healing.
40. A ligand according to claim 38 for use in the treatment of ischemia.
41. A TIE-2 antagonist for use in a method of inhibiting TIE-2 ligand activity in a mammal.

42. An antagonist according to claim 41 which is an antibody capable of specifically binding TIE-2 receptor.

43. An antibody according to claim 42 which is an antibody according to claim 22 or 23.

44. An antagonist according to claim 42 which is a receptorbody according to claim 22.

45. An antagonist according to claim 41 which is a ligand according to claim 15.

46. An antagonist according to any one of claims 41 to 45 for use in a method wherein the mammal is a human.

47. An antagonist according to any one of claims 41 to 46 for use in a method of attenuating or preventing tumor growth in a human.

48. A method of maintaining a TIE-2 receptor expressing cell in culture, which method comprises administering to the TIE-2 receptor expressing cell an effective amount of the ligand of claim 14.

49. A method according to claim 48 wherein the TIE-2 receptor expressing cell is an endothelial cell.

50. A method of identifying a TIE-2 receptor antagonist comprising contacting cells expressing the TIE-2 receptor with: a) a test compound; and b) a ligand according to claim 14 or 15; under conditions permitting binding of the ligand to the receptor and determining

whether the test compound is capable of interfering with the binding of the ligand to the receptor.

51. A polypeptide produced by the method of claim 19.
52. A nucleic acid according to claim 1 or 23, substantially as hereinbefore described with reference to any one of the foregoing Examples.
53. A vector according to claim 6 or 24, substantially as hereinbefore described with reference to any one of the foregoing Examples.
54. A ligand according to any one of claims 12 to 15, substantially as hereinbefore described with reference to any one of the foregoing Examples.
55. A host-vector system according to claim 16, substantially as hereinbefore described with reference to any one of the foregoing Examples.
56. A method according to claim 19, 48 or 50, substantially as hereinbefore described with reference to any one of the foregoing Examples.
57. An antibody according to claim 20 or 33, substantially as hereinbefore described.
58. A receptorbody according to claim 22 or 33, substantially as hereinbefore described with reference to any one of the foregoing Examples.

59. A conjugate according to claim 27 or 33, substantially as hereinbefore described.

60. A composition according to any one of claims 29 to 32 or 33, substantially as hereinbefore described with reference to any one of the foregoing Examples.

61. A TIE-2 antagonist according to claim 41, substantially as hereinbefore described with reference to any one of the foregoing Examples.

62. A ligandbody which specifically binds the TIE-2 receptor or the receptorbody of claim 22, 33 or 58.

63. A ligandbody which comprises a TIE-2 ligand fused to an immunoglobulin constant region.

64. The ligandbody of claim 63 wherein the TIE-2 ligand is TIE-2 ligand according to any one of claims 12 to 15 and the immunoglobulin constant region is the Fc portion of human IgG1.

65. A ligandbody according to any one of claims 62 to 64 for use in a method of treatment of the human or animal body, or in a method of diagnosis.

66. A method of treating a human or animal subject comprising administering to the subject an effective amount of a ligand according to any one of claims 12 to 15, an antibody according to claim 20 or 21, a receptorbody according to claim 22, a conjugate according to claim

27 or 28, a composition according to any one of claims 29 to 32, or a ligandbody according to any one of claims 62 to 65.

67. A method according to claim 66, the method being as defined in any one of claims 36 to 47.

68. A method according to claim 66 further comprising administering a second pharmaceutically active agent.

69. A method according to claim 68 wherein said second pharmaceutically active agent is a cytokine, neurotrophin, interleukin, cytokine antagonist, VEGF, anti-VEGF antibody, VEGF receptorbodies or bFGF.

70. A method of treating leukopenia comprising treating a patient with a therapeutically effective amount of the ligand according to claim 14.

71. A method of treating thrombocytopenia comprising treating a patient with a therapeutically effective amount of the ligand according to claim 14.

72. A method of treating anemia comprising treating a patient with a therapeutically effective amount of the ligand according to claim 14.

73. A method of enhancing bone marrow engraftment during transplantation comprising treating a patient with a therapeutically effective amount of the ligand according to claim 14.

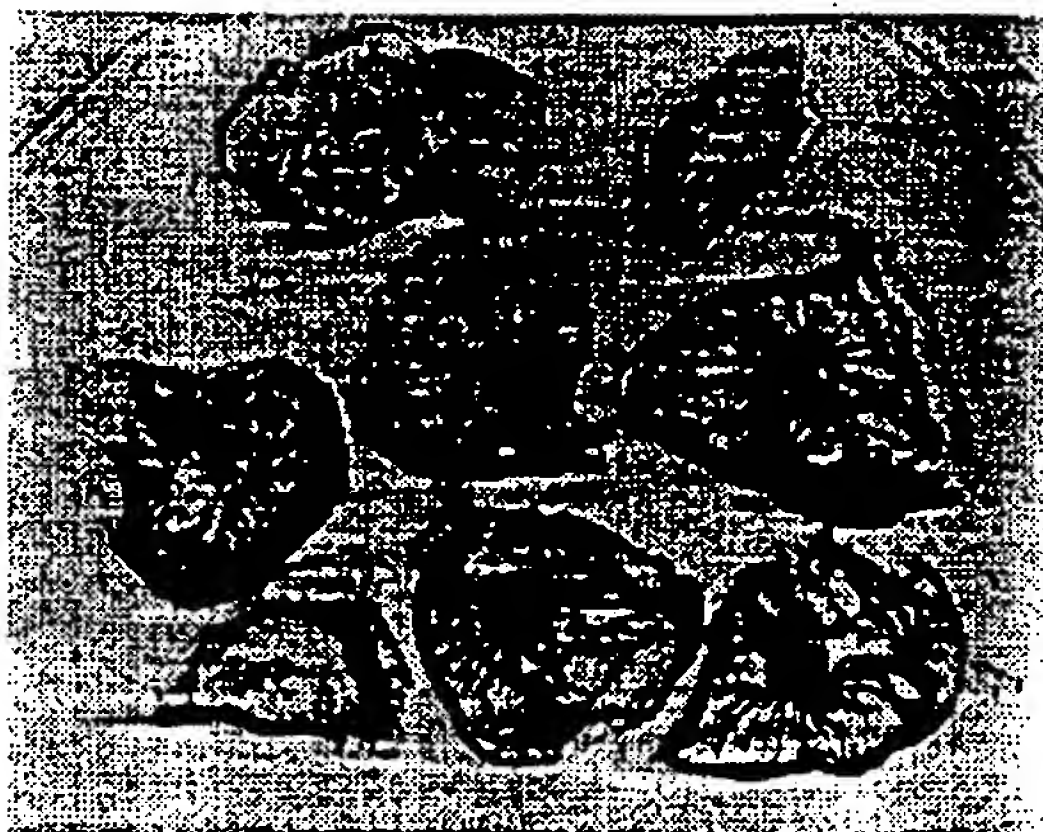
74. A method of treating bone marrow aplasia or myelosuppression caused by radiation, chemical treatment or chemotherapy comprising

treating a patient having such aplasia or myelosuppression with a therapeutically effective amount of the ligand according to claim 14.

75. A method of treating a proliferative disorder of a blood forming organ comprising treating a patient with a therapeutically effective amount of the ligand according to claim 15, an antibody according to claim 20 or 21, a receptorbody according to claim 22, a conjugate according to claim 27 or 28, a composition according to any one of claims 29 to 32, or a ligandbody according to any one of claims 62 to 65.

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Fig.1A.



r EHK-1 ecto/h IgG1 Fc
Gelfoam (6ug)

Fig.1B.



r TIE-2 ecto/h IgG1 Fc
Gelfoam (6ug)

Fig.2.

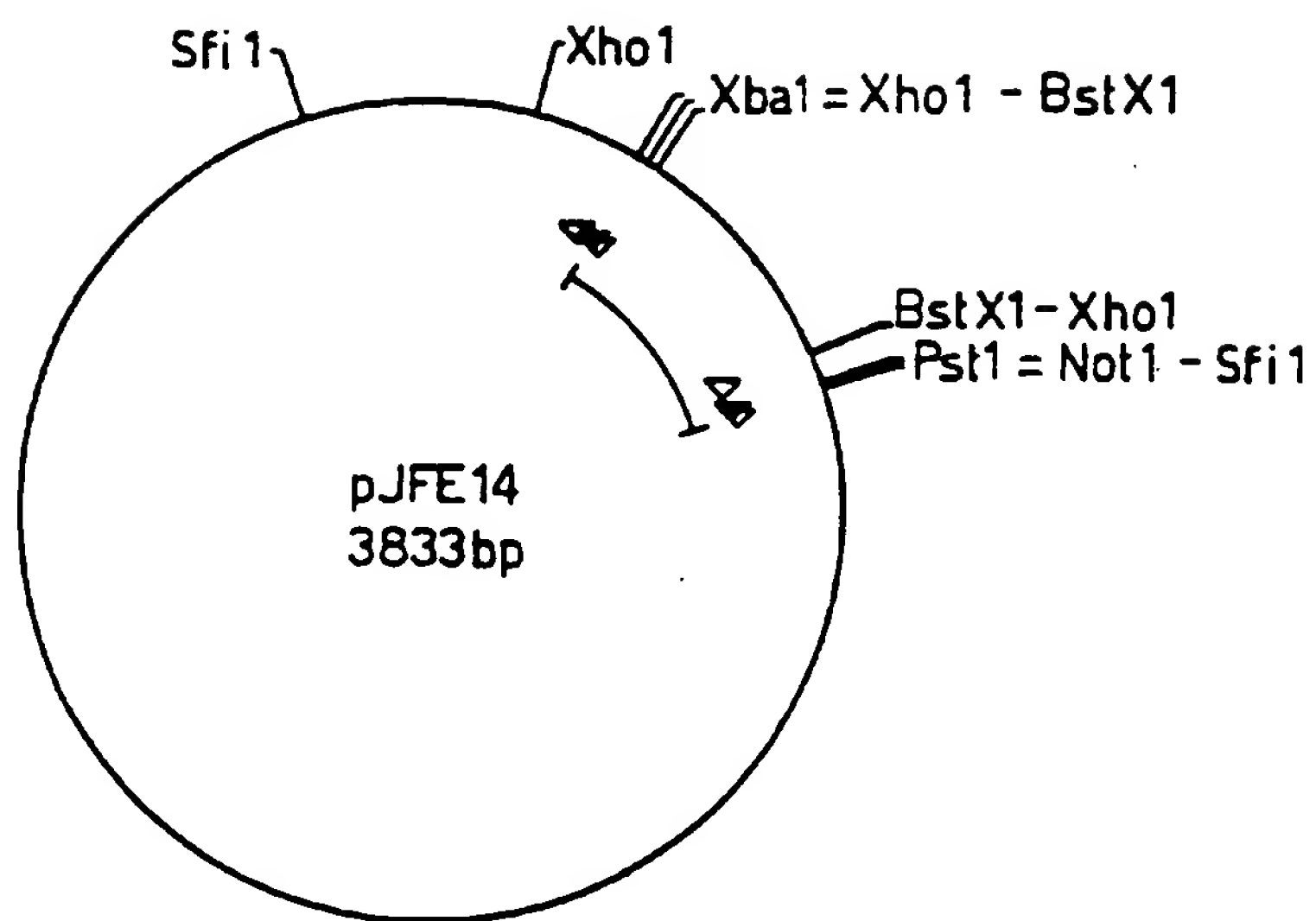


Fig.3.

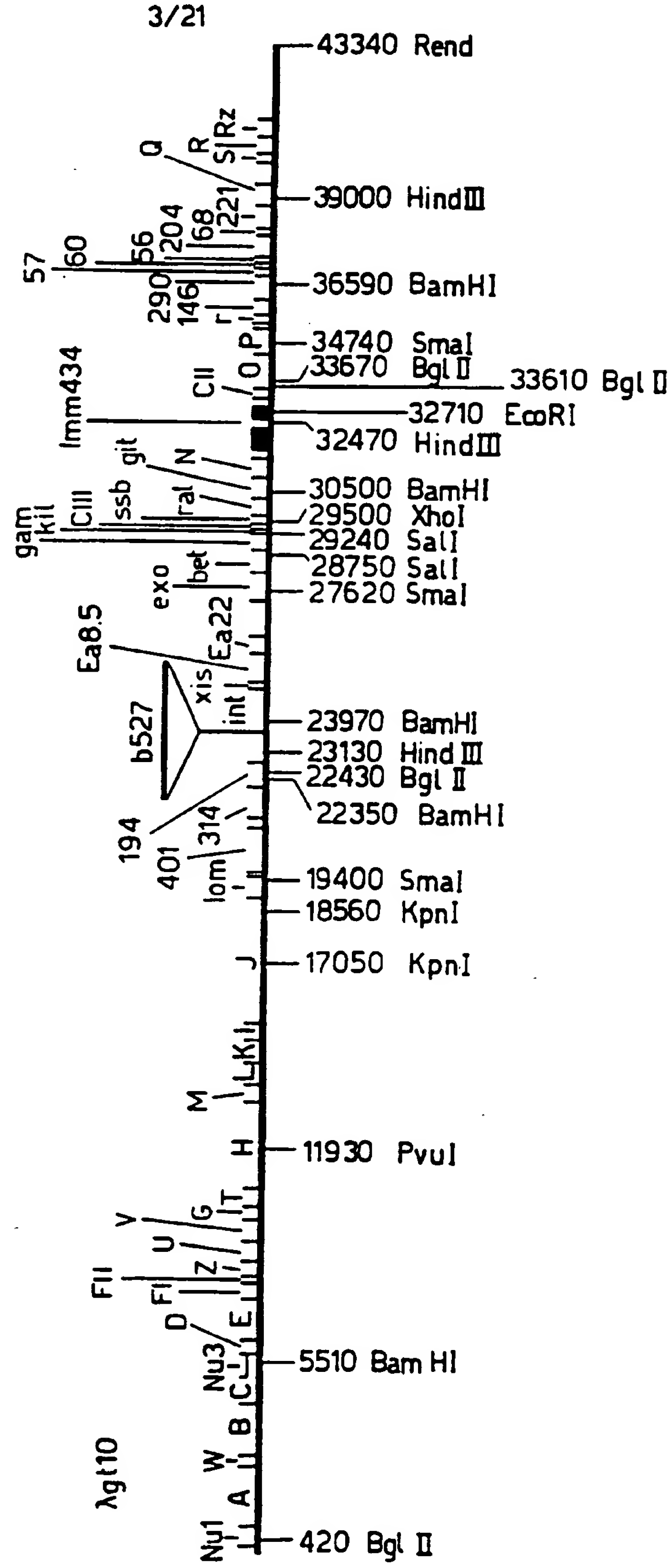


Fig.4.

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10 20 30 40 50 60 70 80
CAGCTGACTCAGCCAGGCTCCATGCTGAACGGTCACACAGAGAGGAAACAATAATTCAGCTACTATGCAATAAATATC

90 100 110 120 130 140 150 160
TCAAGTTTAAAGGAGAAAACATCATTCAGCTGAATATAAAATTTTAAATTTTAAACAAAGCTAACAAATGCTAG

170 180 190 200 210 220 230 240
TTTTCTATGATTTCTTTCAACGGCTTTCTTTGAGGGGAAAGAGTCAACAAACAGCACTTTTACCTGAAATAAGAA

250 260 270 280 290 300 310
CTAGTTTATAGGTCAGAGAAAGGAGCAAGTTTTCGAGAGGCCACCAAGGAGTGTGCTGAGTACA ATG ACA
M T>

320 330 340 350 360 370
GTT TTC CTT TCC TTT GCT TTC CTC GCT GCC ATT CTG ACT CAC ATA GCG TGC ACC AAT CAG
V F L S F A F L A A I L T H : G C S N Q>

380 390 400 410 420 430
CGC CGA AGT CCA GAA AAC AGT GCG AGA AGA TAT AAC CCG ATT CAA CAT GCG CAA TGT GCG
R R S P E N S G R R Y N P I O H G O C A>

440 450 460 470 480 490
TAC ACT TTC ATT CTT CCA GAA CAC GAT GCG AAC TGT CGT GAG AGT ACC ACA GAC CAG TAC
Y T F I L P E H D G N C R E S T T D O Y>

500 510 520 530 540 550
AAC ACA AAC GCT CTG CAG AGA GAT GCT CCA CAC GTG CAA CCG GAT TTC TCT TCC CAG AAA
N T N A L O R D A P H V E P D F S S O K>

560 570 580 590 600 610
CTT CAA CAT CTG GAA CAT GTG ATG GAA AAT TAT ACT CAG TCG CTC CAA AAA CTT GAG AAT
L O H L E H V M E N Y T O W L O K L E N>

620 630 640 650 660 670
TAC ATT GTG GAA AAC ATG AAG TCG GAG ATG GCG CAG ATA CAG CAG AAT GCA GTT CAG AAC
Y I V E N H K S E M A Q I O O N A V Q N>

680 690 700 710 720 730
CAC ACC GCT ACC ATG CTG GAG ATA GGA ACC AGC CTC CTC TCT CAG ACT GCA GAG CAG ACC
H T A T M L E I G T S L L S O T A E O T>

740 750 760 770 780 790
AGA AAG CTG ACA GAT GTT GAG ACC CAG GTA CTA AAT CAA ACT TCT GCA CTT GAG ATA CAG
R K L T D V E T Q V L N Q T S R L E I O>

800 810 820 830 840 850
CTG CTG CAG AAT TCA TTA TCC ACC TAC AAG CTA GAG AAG CAA CTT CTT CAA CAG ACA AAT
L L E N S L S T Y K L E K O L L O O T N>

860 870 880 890 900 910
GAA ATC TTG AAG ATC CAT GAA AAA AAC AGT TTA TTA GAA CAT AAA ATC TTA GAA ATG GAA
E I L K I H E K N S L L E H K I L E H E>

920 930 940 950 960 970
CGA AAA CAC AAG GAA GAG TTG GAC ACC TTA AAG GAA GAG AAA GAG AAC CTT CAA GCG TTC
G K H K E E L D T L K E E K E N L O G L>

980 990 1000 1010 1020 1030
GTT ACT CGT CAA ACA TAT ATA ATC CAG GAG CTC GAA AAG CAA TTA AAC ACA GCT ACC ACC
V T R C T Y I I O E L E K O L N R A T T>

1040 1050 1060 1070 1080 1090
AAC AAC AGT GTC CTT CAG AAG CAG CAA CTG GAG CTG ATG CAC ACA GTC CAC AAC CTT GTC
N N S V L O K O O L E L M D T V H N L V>

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Fig.4. (Cont.)

1100 1110 1120 1130 1140 1150
AAT CTT TGC ACT AAA GAA GGT GTT TTA CTA AAG GGA GGA AAA AGA GAG GAA GAG AAA CCA
N L C T K E G V L L K G G K R E E E K P>
1160 1170 1180 1190 1200 1210
TTT AGA GAC TGT GCA GAT GTA TAT CAA GGT GGT TTT AAT AAA AGT GGA ATC TAC ACT ATT
F R D C A D V Y O A C F N K S G I Y T I>
1220 1230 1240 1250 1260 1270
TAT ATT AAT AAT ATG CCA GAA CCC AAA AAG GTG TTT TGC AAT ATG GAT GTC AAT GCG GCA
Y I N N N P E P K K V F C N M D V N G G>
1280 1290 1300 1310 1320 1330
GGT TGC ACT GTA ATA CAA CAT CGT GAA GAT GGA AGT CTA GAT TTC CAA AGA GCG TCG AAG
G W T V I Q H R E D G S L D F Q R G W K>
1340 1350 1360 1370 1380 1390
GAA TAT AAA ATG GGT TTT GGA AAT CCC TCC GGT GAA TAT TCG CTC GCG AAT GAG TTT ATT
E Y K M G F G N P S G E Y W L G N E F I>
1400 1410 1420 1430 1440 1450
TTT GCG ATT ACC AGT CAG AGG CAG TAC ATG CTA AGA ATT GAG TTA ATG GAC TCG GAA GCG
F A I T S Q R Q Y M L R I E L M D W E G>
1460 1470 1480 1490 1500 1510
AAC CGA GCG TAT TCA CAG TAT GAC AGA TTC CAC ATA GGA AAT GAA AAG CAA AAC TAT AGC
N R A Y S Q Y D R F H I G N E K Q N Y R>
1520 1530 1540 1550 1560 1570
TTG TAT TTA AAA GGT CAC ACT GCG ACA GCA GGA AAA CAG AGC AGC CTG ATC TTA CAC GGT
L Y L K G H T G T A G K Q S S L I L H G>
1580 1590 1600 1610 1620 1630
CGT GAT TTC AGC ACT AAA GAT CGT GAT AAT GAC AAC TGT ATG TCG AAA TGT GCG CTC ATG
A D F S T K D A D N D N C M C K C A L M>
1640 1650 1660 1670 1680 1690
TTA ACA GGA GGA TCG TCG TTT GAT CGT TGT GCG CCC TCC AAT CTA AAT GGA ATG TTC TAT
L T G G W W F D A C G P S N L N G M F Y>
1700 1710 1720 1730 1740 1750
ACT GCG GGA CAA AAC CAT GGA AAA CTG AAT GCG ATA AAG TCG CAC TAC TTC AAA GCG CCC
T A G O N H G K L M G I K W H Y F K G P>
1760 1770 1780 1790 1800 1810
AGT TAC TCC TTA CGT TCC ACA ACT ATG ATG ATT GGA CGT TTA GAT TTT TGA AAG CGCAATGT
S Y S L R S T T M M I R P L D F
1820 1830 1840 1850 1860 1870 1880 1890
CAGAACGATTATGAAAGCAAGAAATCCGAGAGCTCCAGGTGAGAACTGTTTGAAACTTCAGAACCAACA
1900 1910 1920 1930 1940 1950 1960 1970
ATATGCTCTCCCTTCCAGCAATAAGTCTAGTTATGTGAAGTCAACAGGTTCTTGACCGTGAACTCGAGCCGTTTGAC
1980 1990 2000 2010 2020 2030 2040 2050
TTCACAGAGTCTCTACTTCCGCTGACAGTCTCAGTCCGCTGACTATAGAAAACCTCCACTGACTCTCCGCTTTAAAA
2060 2070 2080 2090 2100 2110 2120 2130
ACCGAAGAACTGCTGAGCTTCTGCTCTCAAACTACTACTGACCTTATTTTGGAACTATGTTAGCCAGATCATTAAT
2140
ATGGTAATTTC

Fig.5.

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10 20 30 40 50 60 70 80
CACTGACTCAGCCAGCTCCATGCTGACCGTCCACAGAGAGGAAACATTAATCTCAGCTACTATGCAATAATATC
90 100 110 120 130 140 150 160
TCAAGTTTAAAGGAGAAAACATCATTGCACTGAATAAAAAATTTAAATTTTAGAACAAAGCTAACAAATGCTAG
170 180 190 200 210 220 230 240
TTTCTATGATTCTTCTTCAAGGCTTTCTTTGAGGCGAAGAGTCAACAAACAAGCAGTTTACTGAAATTAAGAA
250 260 270 280 290 300 310
CTAGTTTAAAGGTCAGAACAAAGGAGCAAGTTTTCGAGAGGACAGGAGGAGTGTCTGCGAGTACA ATG ACA
H T>
320 330 340 350 360 370
GTT TTC CTT TCC TTT OCT TTC CTC GCT GGC ATT CTG ACT CAC ATA GCG TGC AGC AAT CAG
V F L S F A F L A A I L T H I G C S N Q>
380 390 400 410 420 430
CCG CCA AGT CCA GAA AAC AGT GCG AGA AGA TAT AAC GCG ATT CAA CAT GCG CAA TGT GCG
R R S P E N S G R R V N R I O H G O C A>
440 450 460 470 480 490
TAC ACT TTC ATT CTT CCA GAA CAC GAT GCG AAC TGT GGT GAG AGT AGC ACA GAC CAG TAC
Y T F I L P E H D G N C R E S T T D Q Y>
500 510 520 530 540 550
AAC ACA AAC GCT CTG CAG AGA GAT GCT CCA CAC CTG GAA CCG GAT TTC TCT TCC CAG AAA
N T N A L Q R D A P H V E P D F S S O K>
560 570 580 590 600 610
CTT CAA CAT CTG GAA CAT GTG ATG GAA AAT TAT ACT CAG TCG CTG CAA AAA CTT CAG AAT
L Q H L E H V H E N Y T Q W L O K L E N>
620 630 640 650 660 670
TAC ATT GTG GAA AAC ATG AAG TCG GAG ATG GCG CAG ATA CAG CAG AAT GCA GTT CAG AAC
Y I V E N H K S E H A Q I Q O N A V O N>
680 690 700 710 720 730
CAC ACC GCT ACC ATG CTG GAG ATA GGA ACC ACC CTC CTC TCT CAG ACT GCA GAG CAG ACC
H T A T H L E I G T S L L S O T A E O T>
740 750 760 770 780 790
AGA AAG CTG ACA GAT GTT GAG ACC CAG GTA CTA AAT CAA ACT TCT GCA CTT GAG ATA CAG
R K L T D V E T Q V L N Q T S R L E I Q>
800 810 820 830 840 850
CTG CTG GAG AAT TCA TTA TCC ACC TAC AAG CTA GAG AAG CAA CTT CTT CAA CAG ACA AAT
L L E N S L S T Y K L E K O L L O O T N>
860 870 880 890 900 910
GAA ATC TTG AAG ATC CAT GAA AAA AAC AGT TTA TTA GAA CAT AAA ATC TTA GAA ATG GAA
E I L K I H E K N S L L E H K I L E H E>
920 930 940 950 960 970
GCA AAA CAC AAG GAA GAG TTG CAC ACC TTA AAG GAA CAG AAA CAG AAC CTT CAA GCG TTG
G K H K E E L D T L K E E K E N L O G L>
980 990 1000 1010 1020 1030
GTT ACT GGT CAA ACA TAT ATA ATC CAG CAG CTC GAA AAG CAA TTA AAC AGA GGT ACC ACC
V T R O T Y I I O E L E K O L N R A T T>
1040 1050 1060 1070 1080 1090
AAC AAC AGT GTC CTT CAG AAG CAG CAA CTC CAG CTC ATG GAC ACA GTC CAC AAC CTT GTC
N N S V L Q K O O L E L H D T V H N L V>

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Fig.5. (Cont.)

1100 1110 1120 1130 1140 1150
AAT CTT TGC ACT AAA GAA GTT TTA CTA AAG CGA CGA AAA AGA GAG GAA GAG AAA CCA TTT
N L C T K E V L L K G G K R E E E K P F>
1160 1170 1180 1190 1200 1210
AGA GAC TGT CGA GAT GTA TAT CAA GGT GGT TTT AAT AAA AGT CGA ATC TAC ACT ATT TAT
R D C A D V Y Q A G F N K S G I Y T I Y>
1220 1230 1240 1250 1260 1270
ATT AAT AAT ATG CGA GAA CCC AAA AAG GTG TTT TGC AAT ATG GAT CTC AAT GCG CGA GGT
I H H H P E P K K V F C N H D V N G G G>
1280 1290 1300 1310 1320 1330
TGG ACT GTA ATA CAA CAT GGT GAA GAT CGA AGT CTA GAT TTC CAA AGA GCG TGC AAG GAA
W T V I Q H R E D G S L D F Q R G W K E>
1340 1350 1360 1370 1380 1390
TAT AAA ATG GGT TTT GGA AAT CCC TCC GGT GAA TAT TGC CTC GCG AAT GAG TTT ATT TTT
Y K H G F C N P S G E Y W L G N E F I F>
1400 1410 1420 1430 1440 1450
GCG ATT ACC AGT CAG AGC CAG TAC ATG CTA AGA ATT GAG TTA ATG GAC TGC GAA GCG AAC
A I T S Q R O Y H L R I E L M D W E G N>
1460 1470 1480 1490 1500 1510
CGA GCG TAT TCA CAG TAT CAC AGA TTC CAC ATA CGA AAT GAA AAG CAA AAC TAT AGC TTG
R A Y S Q Y D R F H I G H E K Q N Y R L>
1520 1530 1540 1550 1560 1570
TAT TTA AAA GGT CAC ACT GCG ACA CGA CGA AAA CAG AGC AGC CTC ATC TTA CAC GGT GGT
Y L K G H T C T A G K Q S S L I L H G A>
1580 1590 1600 1610 1620 1630
GAT TTC ACC ACT AAA GAT GGT GAT AAT GAC AAC TGT ATG TCC AAA TGT GCG CTC ATG TTA
D F S T K D A D N D N C H C K C A L H L>
1640 1650 1660 1670 1680 1690
ACA CGA CGA TGC TGC TTT GAT GGT TGT GCG CCC TCC AAT CTA AAT CGA ATC TTC TAT ACT
T C G C W W F D A C G P S H L H G H F Y T>
1700 1710 1720 1730 1740 1750
GCG CGA CAA AAC CAT CGA AAA CTC AAT GCG ATA AAG TCG CAC TAC TTC AAA GCG CCC AGT
A G O N H G K L N G I K W H Y F K G P S>
1760 1770 1780 1790 1800 1810
TAC TCC TTA GGT TCC ACA ACT ATC ATG ATT CGA CCT TTA GAT TTT TGA AAGCCCAATGTCAGAA
Y S L R S T T H H I R P L D F>
1820 1830 1840 1850 1860 1870 1880 1890
CCGATTATCAAGCAACAAAGAAATCCGACAGCTCCAGGTGAGAACTGTTGAAAATTCAGAACCAACAAATATT
1900 1910 1920 1930 1940 1950 1960 1970
GTCTCCCTTCAGCAATAAGTGTAGTTATGTAAGTCACCAAGCTTCTTGACCTCAATCTGAGCTGTTGACTTCAC
1980 1990 2000 2010 2020 2030 2040 2050
AAGAGTCTCTACTTCGGGTGACACTGCTCAGCTGCTCGACTATAGAAAATCCACTGCTGCTGGCTTTAAAAAGCGA
2060 2070 2080 2090 2100 2110 2120 2130
AGAAACTGCTGAGCTTCTGCTGCTTCACTACTGACCTTATTTTGAAGTATGCTAGCCAGATCATATATATGCT
2140
TAATTC

Fig.6. 8/21

10 20 30 40 50 60 70 80
GAATTCCTGGGTGGTGTATCTCTCCAGCCTTCAGGGAGGGGAACAACACTGTAGCATCTGGGAGAGAGGAACAAA
90 100 110 120 130 140 150 160
GGACCGTGAAAGCTGCTCTGTAAAGCTGACACAGCCCTCCCAAGTGAGCAGGACTGTTCTTCCCACTGCAATCTGACAG
170 180 190 200 210 220 230 240
TTTACTGCATGCTGGAGAGAACACAGCAGTAAAAACCAAGTTTCTACTGGAAAAGAGGAAGAGAAAGACTTTCATTG
250 260 270 280 290 300 310 320
ACGGACCCAGCCATGCCAGCGTAGCAGCCCTGCGTTTCAGACGGCAGCAGCTCGGGACTCTGGAGCTGTGTTTGCCTCA
330 340 350 360 370 380
AGTTTGTAAAGCTGCTGTTTATTACTGAAGAAAGA ATG TGG CAG ATT GTT TTC TTT ACT CTG AGC TGT
H W Q I V F F T L S C>
390 400 410 420 430 440
GAT CTT GTC TTG GCC GCA GCC TAT AAC AAC TTT CCG AAG AGC ATG GAC AGC ATA GGA AAG
D L V L A A A Y N N F R K S H D S I G S>
450 460 470 480 490 500
AAG CAA TAT CAG GTC CAG CAT GGG TCC TGC AGC TAC ACT TTC CTC CTC CCA GAG ATG GAC
K Q Y Q V Q H G S C S Y T F L L P E H D>
510 520 530 540 550 560
AAC TGC CGC TCT TCC TCC AGC CCC TAC GTG TCC AAT GCT GTG CAG AGG GAC GCG CCG CTC
N C R S S S S P Y V S N A V Q R D A P L>
570 580 590 600 610 620
GAA TAC GAT GAC TCG GTG CAG AGG CTG CAA GTG CTG GAG AAC ATC ATG GAA AAC AAC ACT
E S D D S V Q R L Q V L E N I H E K H T>
630 640 650 660 670 680
CAG TGC CTA ATG AAG CTT GAG AAT TAT ATC CAG GAC AAC ATG AAG AAA GAA ATG GTA GAG
Q K L H K L E N Y I Q D N H K K E H V E>
690 700 710 720 730 740
ATA CAG CAG AAT GCA GTA CAG AAC CAG ACG GCT GTG ATG ATA GAA ATA GCG ACA AAC CTC
I C Q N A V Q N Q T A V H I E I G T N L>
750 760 770 780 790 800
TTG AAC CAA ACA GCT GAG CAA ACG CCG AAG TTA ACT GAT GTG GAA GCC CAA GTA TTA AAT
L N Q T A E Q T R K L T D V E A Q V L N>
810 820 830 840 850 860
CAG ACC ACC AGA CTT GAA CTT CAG CTC TTG GAA CAC TCC CTC TCG ACA AAC AAA TTG GAA
Q T T R L E L Q L L E H S L S T N K L E>
870 880 890 900 910 920
AAA CAG ATT TTG GAC CAG ACC AGT GAA ATA AAC AAA TTG CAA GAT AAG AAC AGT TTC CTA
K C I L D Q T S E I N K L Q D K N S F L>
930 940 950 960 970 980
GAA AAG AAG GTG CTA GCT ATG GAA GAC AAG CAC ATC ATC CAA CTA CAG TCA ATA AAA GAA
E K K V L A H E D K H I I Q L Q S I K E>
990 1000 1010 1020 1030 1040
GAG AAA GAT CAG CTA CAG GTG TTA GTA TCC AAG CAA AAT TCC ATC ATT GAA GAA CTA GAA
E N D Q L Q V L V S K Q N S I I E E L E>
1050 1060 1070 1080 1090 1100
AAA AAA ATA GTG ACT GCC ACG GTG AAT AAT TCA GTT CTT CAA AAG CAG CAA CAT GAT CTC
K E S V T A T V N N S V L C K Q Q H D L>

Fig.6. (Cont.) 9/21

1110 1120 1130 1140 1150 1160
ATG GAG ACA GTT AAT AAC TTA CTG ACT ATG TCC ACA TCA AAC TCA GCT AAG GAC CCC
H E T V N N L L T H M S T S N S A K D P>
1170 1180 1190 1200 1210 1220
ACT GTT OCT AAA GAA GAA CAA ATC AGC TTC AGA GAC TGT GCT GAA GTA TTC AAA TCA GGA
T V A K E E O I S F R D C A E V F K S G>
1230 1240 1250 1260 1270 1280
CAC ACC ACA AAT GGC ATC TAC AGC TTA ACA TTC CCT AAT TCT ACA GAA GAG ATC AAG GCC
H T T N G I Y T L T F P N S T E E I K A>
1290 1300 1310 1320 1330 1340
TAC TGT GAC ATG GAA GCT GGA GGA GGC GGC TGG ACA ATT ATT CAG CGA CGT GAG GAT GGC
Y C D H E A G G G G W T I I O R R E D G>
1350 1360 1370 1380 1390 1400
AGC GTT GAT TTT CAG AGG ACT TGG AAA GAA TAT AAA GTG GGA TTT GGT AAC CCT TCA GGA
S V D F Q R T W K E Y K V G F G N P S G>
1410 1420 1430 1440 1450 1460
GAA TAT TGG CTG GGA AAT CAG TTT GTT TCG CAA CTG ACT AAT CAG CAA CGC TAT CTG CTT
E Y W L G N E F V S Q L T N O Q R Y V L>
1470 1480 1490 1500 1510 1520
AAA ATA CAC CTT AAA GAC TGG GAA GGG AAT GAG GCT TAC TCA TTG TAT GAA CAT TTC TAT
K I H L K D W E G N E A Y S L Y E H F Y>
1530 1540 1550 1560 1570 1580
CTC TCA AGT GAA GAA CTC AAT TAT AGG ATT CAC CTT AAA GGA CTT ACA GGG ACA GCC GGC
L S S E E L N Y R I H L K G L T G T A G>
1590 1600 1610 1620 1630 1640
AAA ATA AGC AGC ATC AGC CAA CCA GGA AAT GAT TTT AGC ACA AAG GAT GGA GAC AAC GAC
K I S S I S Q P G N D F S T K D G D N D>
1650 1660 1670 1680 1690 1700
AAA TGT ATT TGC AAA TGT TCA CAA ATG CTA ACA GGA GGC TGG TGG TTT GAT GCA TGT GGT
K C I C K C S Q H L T G G W W F D A C G>
1710 1720 1730 1740 1750 1760
CCT TCC AAC TTC AAC GGA ATG TAC TAT CCA CAG AGG CAG AAC ACA AAT AAG TTC AAC GGC
P S N L N G H Y Y P Q R Q N T N K F N G>
1770 1780 1790 1800 1810 1820
ATT AAA TGG TAC TAC TGG AAA GGC TCA GGC TAT TGG CTC AAG GCC ACA ACC ATG ATG ATC
I K W Y Y W K G S G Y S L K A T T H H I>
1830 1840 1850 1860 1870 1880 1890 1900
CGA CCA GCA GAT TTC TAAACATCCCACTCCACCTGAGGAAGTGTCTCGAACTATTTTCAAAGACTTAAGCCCACT
R P A D F>
1910 1920 1930 1940 1950 1960 1970 1980
GCACTGAAAGTCACCGCTGCCCACTGTGTCTCTTTCCACCAAGAGGGGGCTGTCTCGCTGACGGGACCCACATGCT
1990 2000 2010 2020 2030 2040 2050 2060
CCAGATTAGAGCCTGTAAACTTTATCACTTAAACTTGCATCACTTAACGGACCAAGCAAGACCCTAAACATCCATAAT
2070 2080 2090 2100 2110 2120 2130 2140
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2150 2160 2170 2180 2190 2200 2210 2220
CCAAGAATGTTATGTGCAAGTTTATCACTAAATAACTGCAAAACAGAACTTATGTTATACAATACAGATCATCTTGA
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Fig.7.

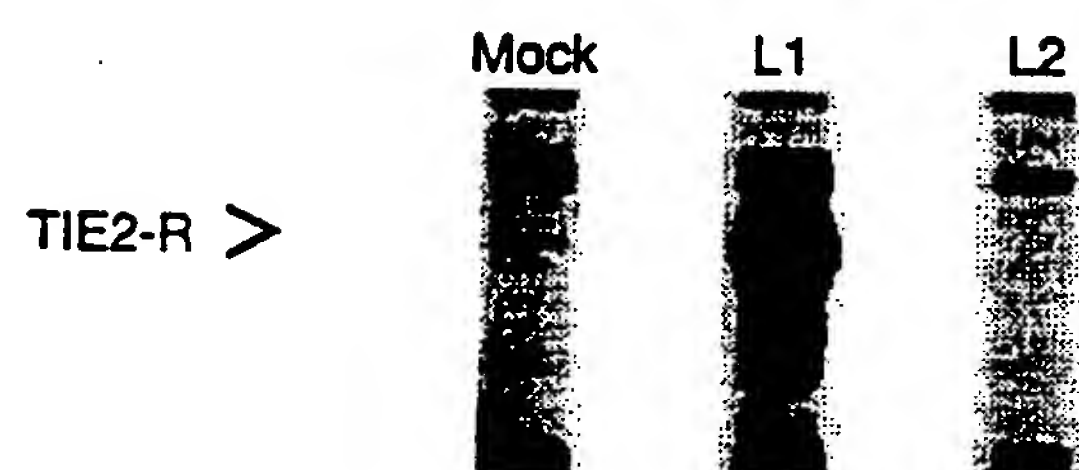
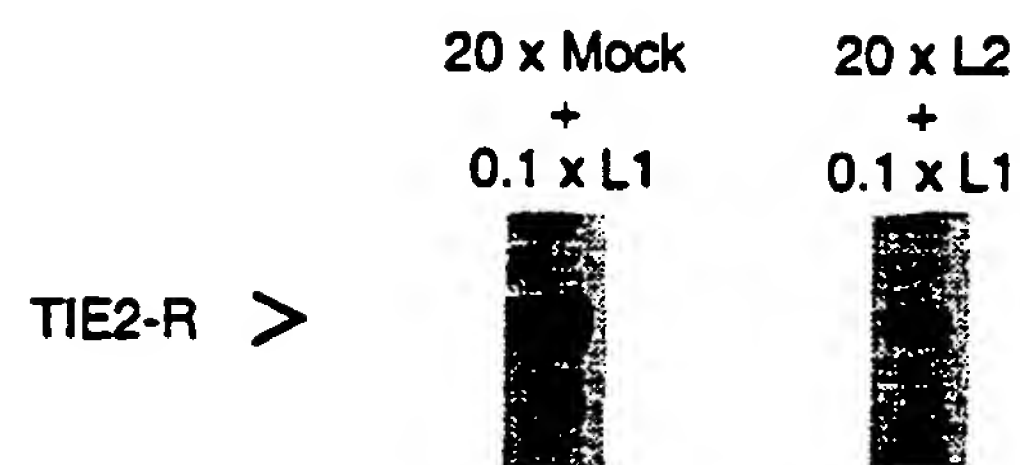
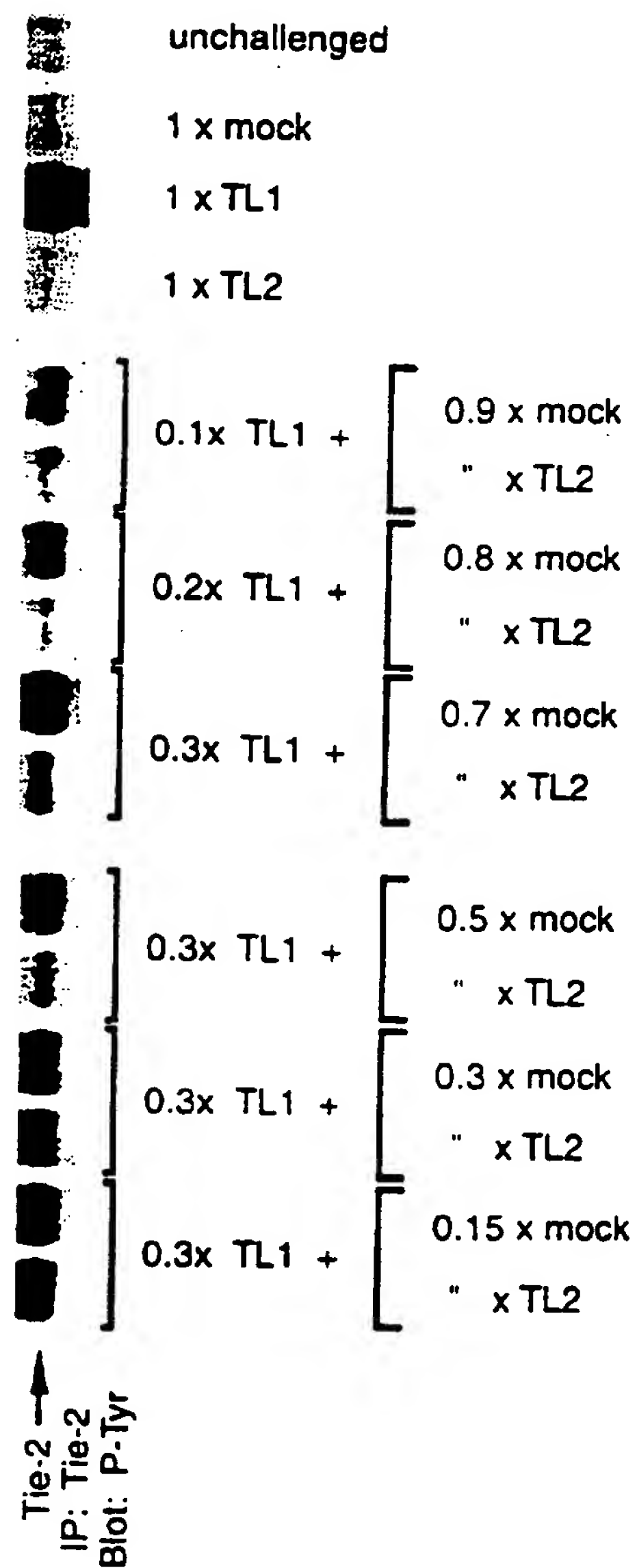


Fig.8.



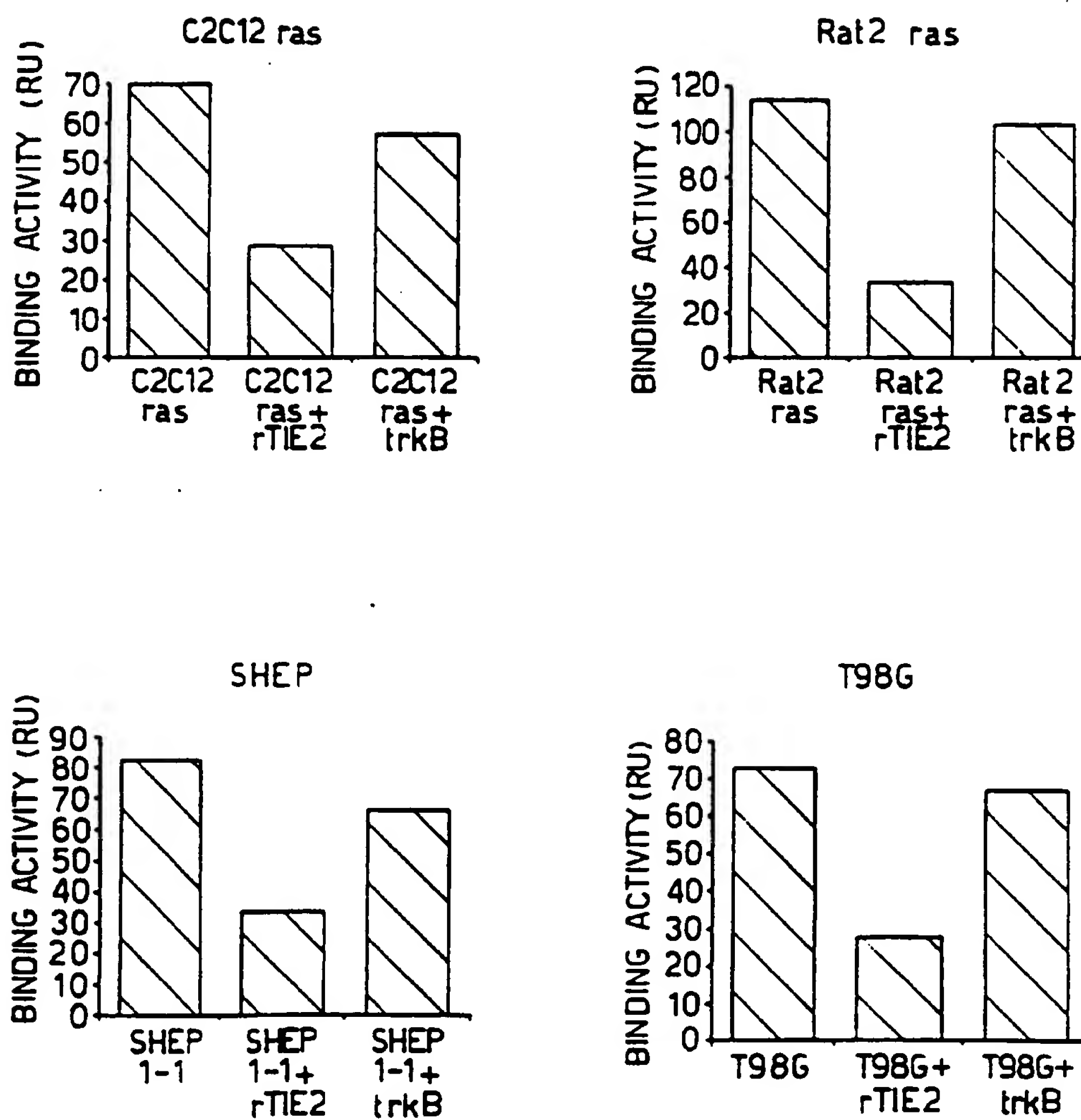
11/21

Fig.9.



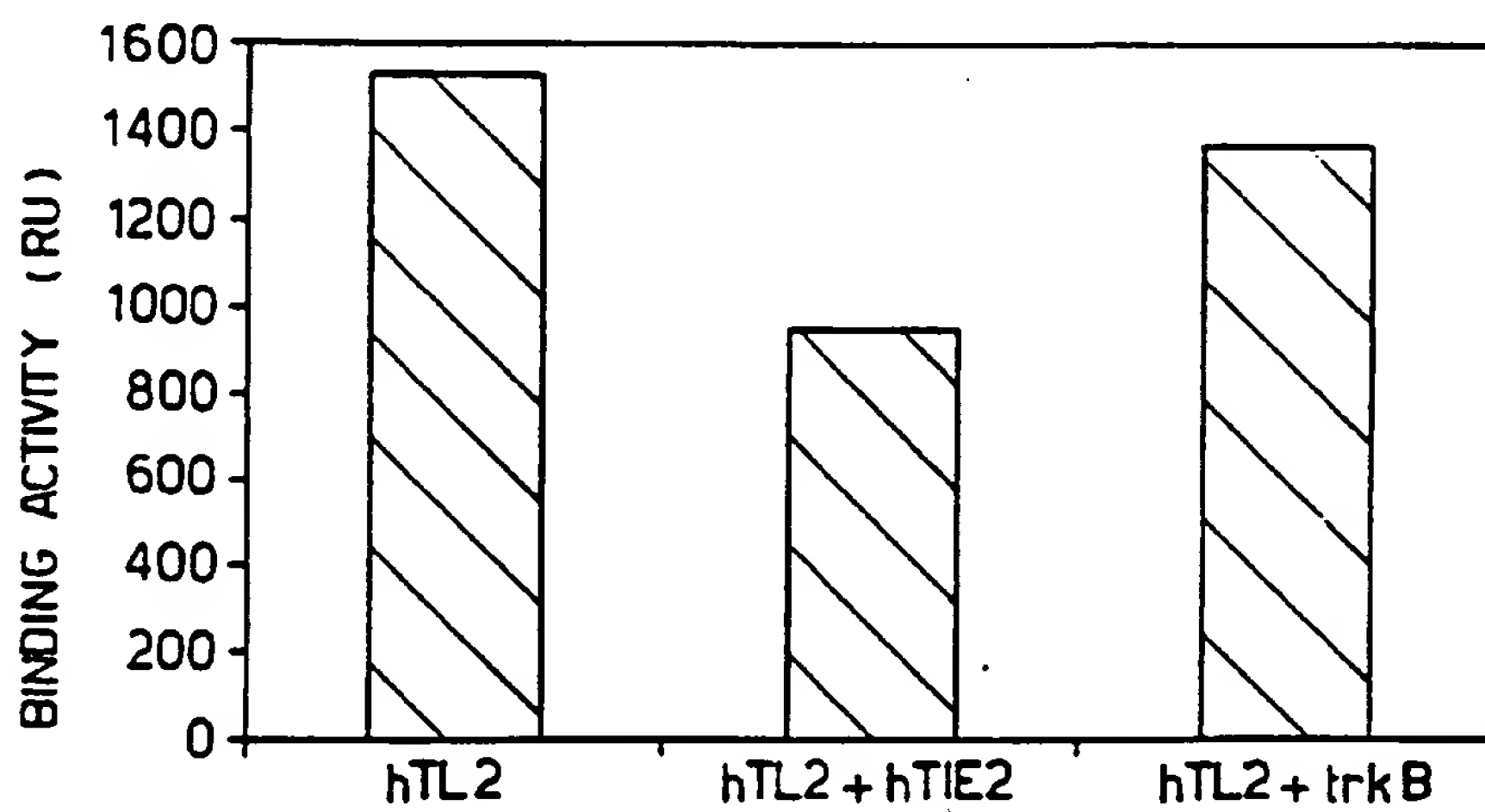
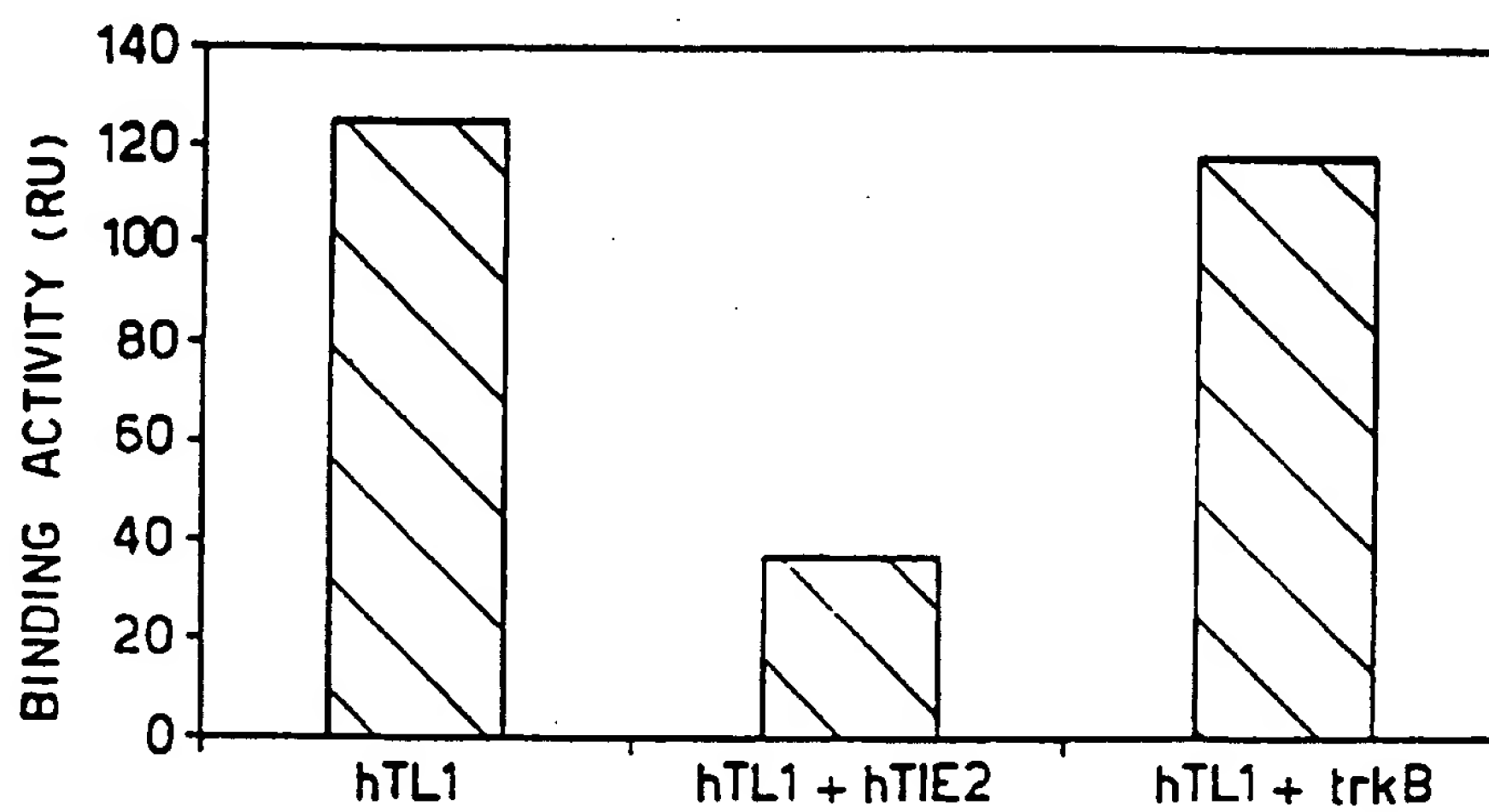
12/21

Fig.10.



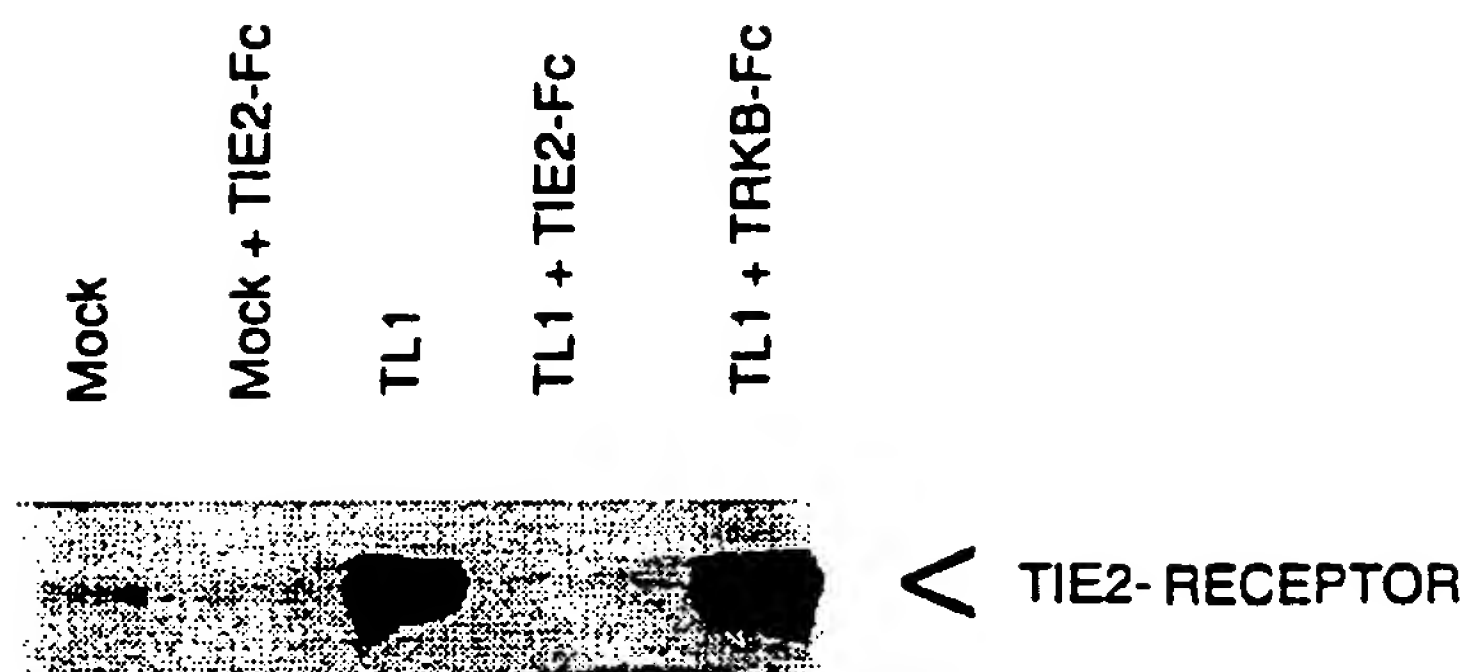
13/21

Fig.11.



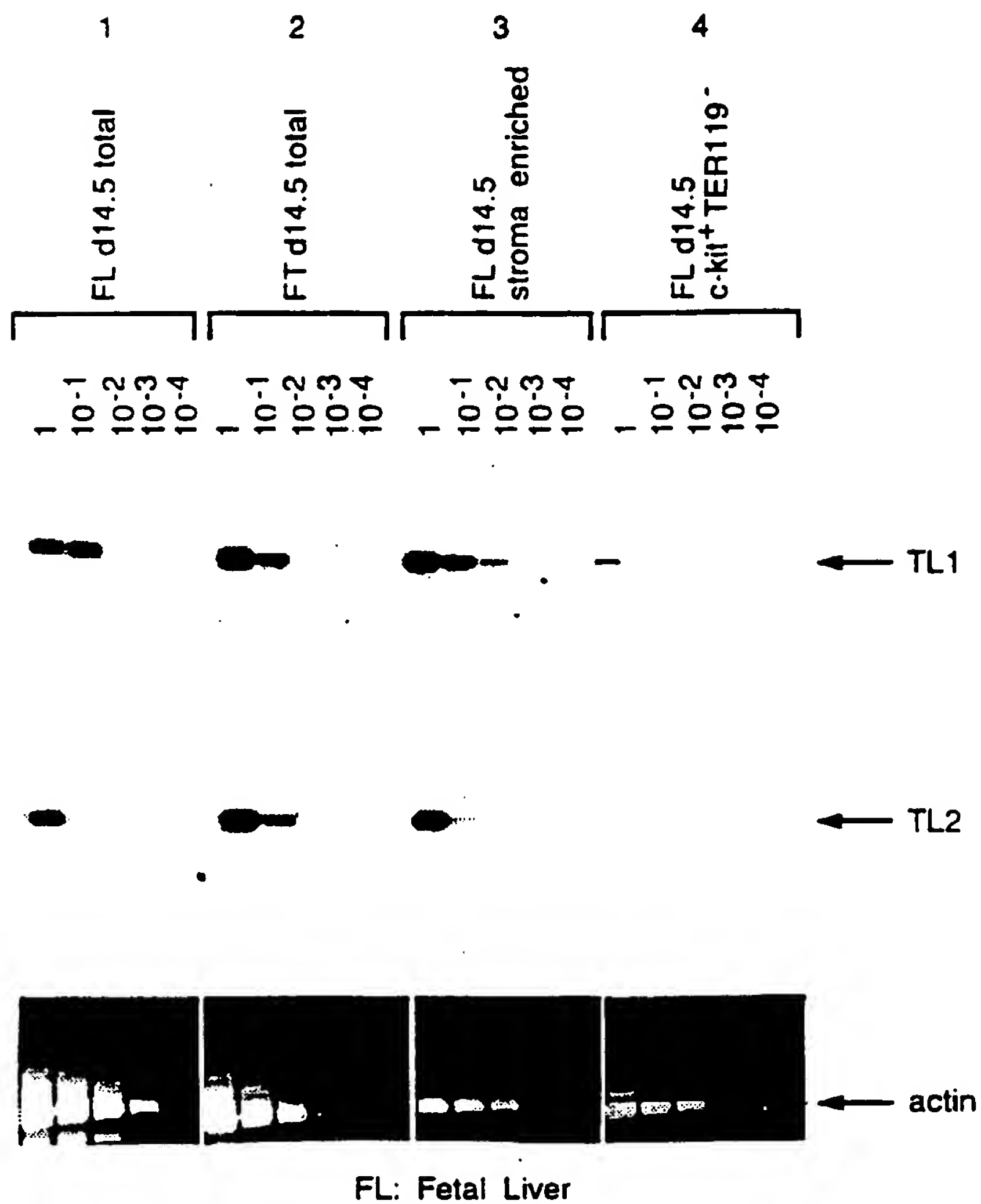
14/21

Fig.12.



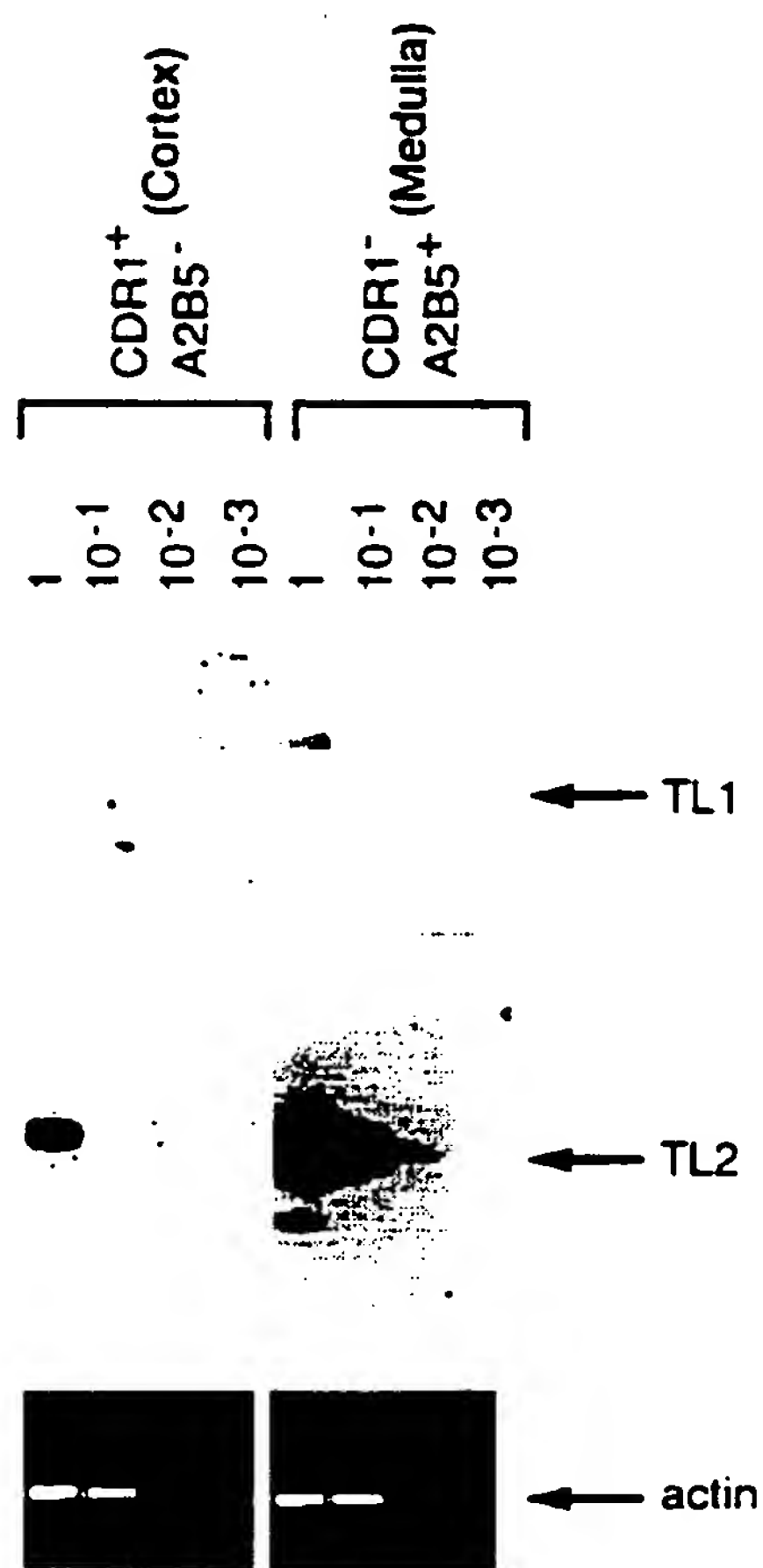
15/21

Fig.13.



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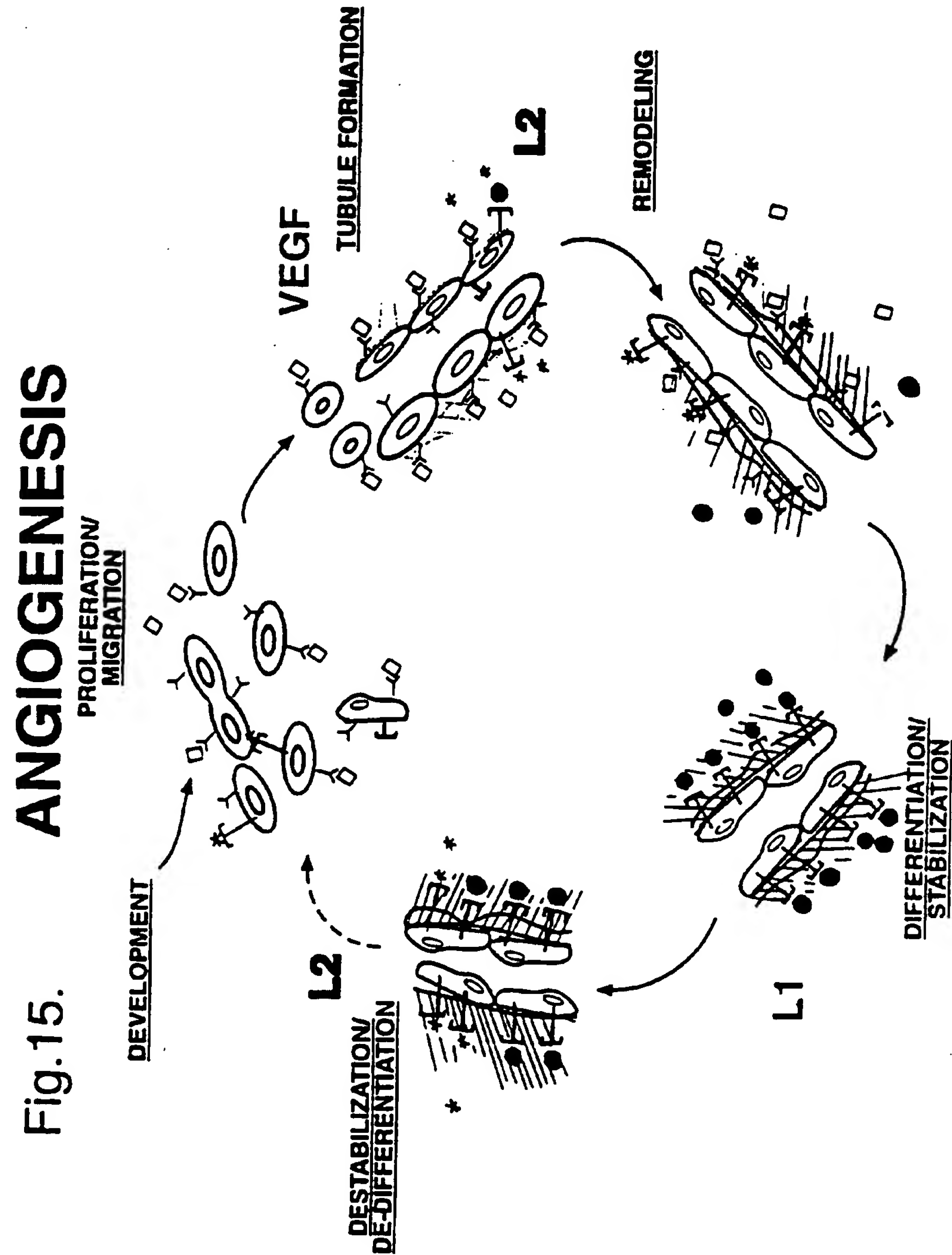
Fig. 14.



Fetal Thymus E17.5

CDR1⁺ : Cortical stromal cellsA2B5⁺ : Medulla stromal cells

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Fig.16.

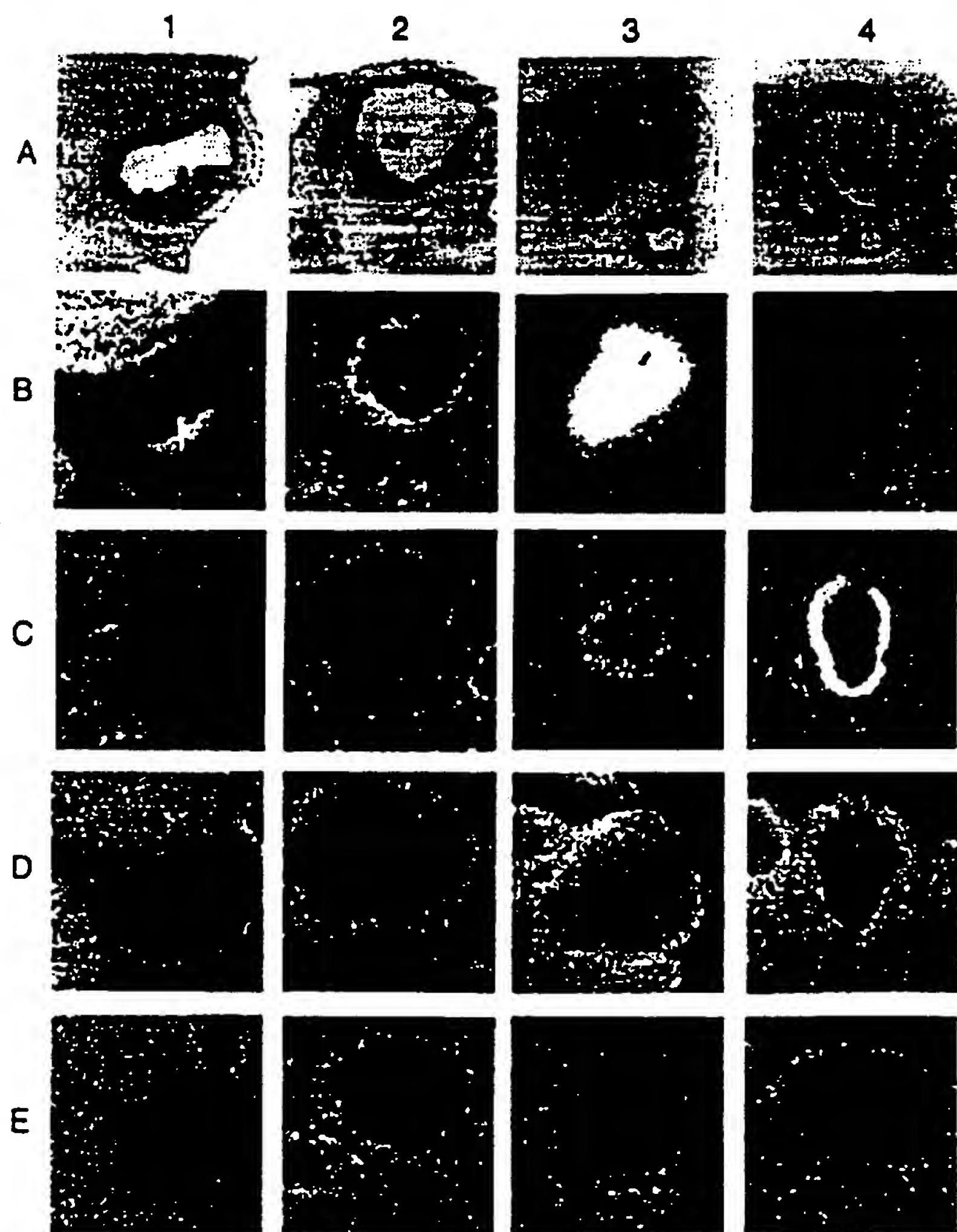
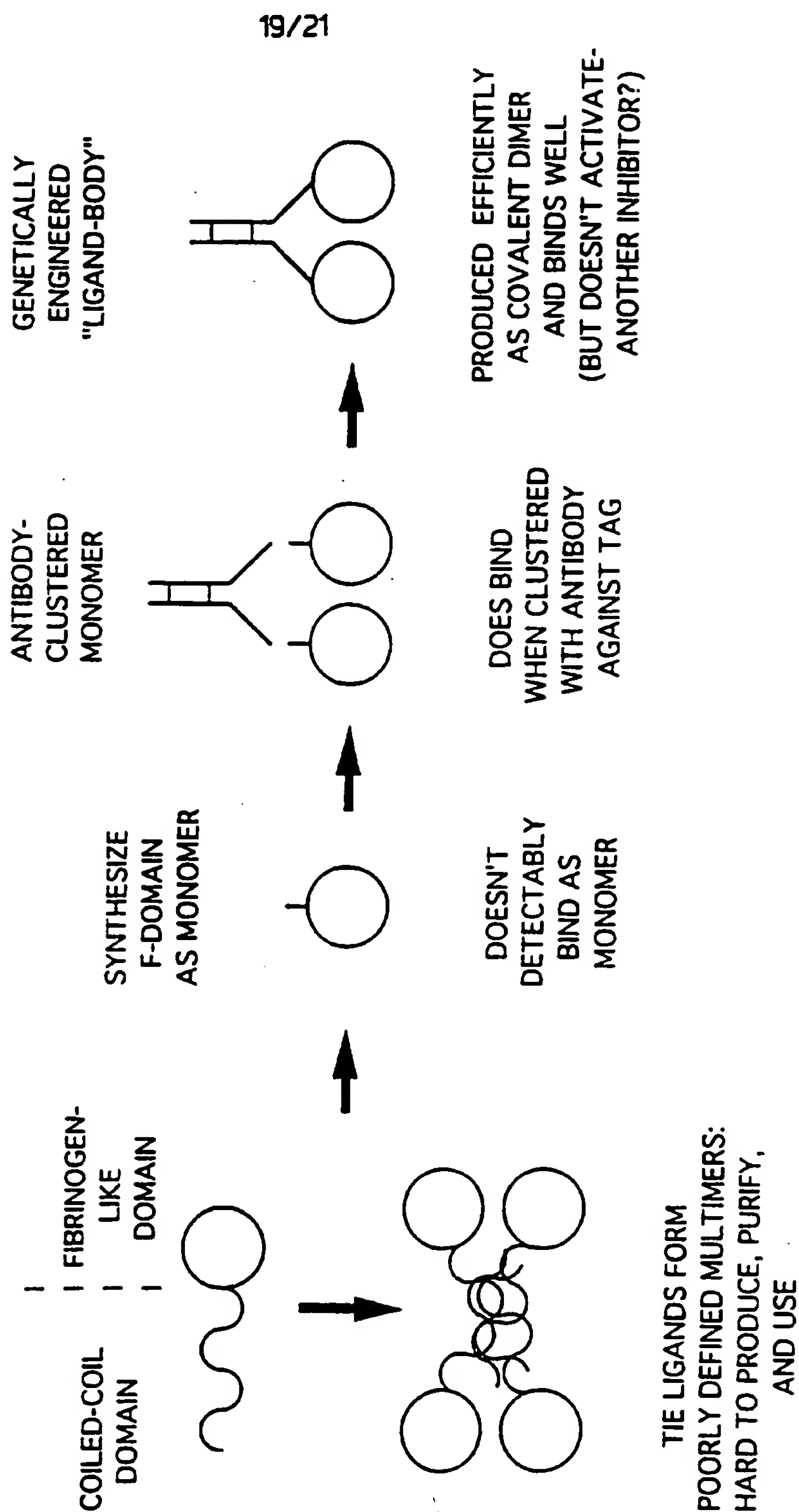


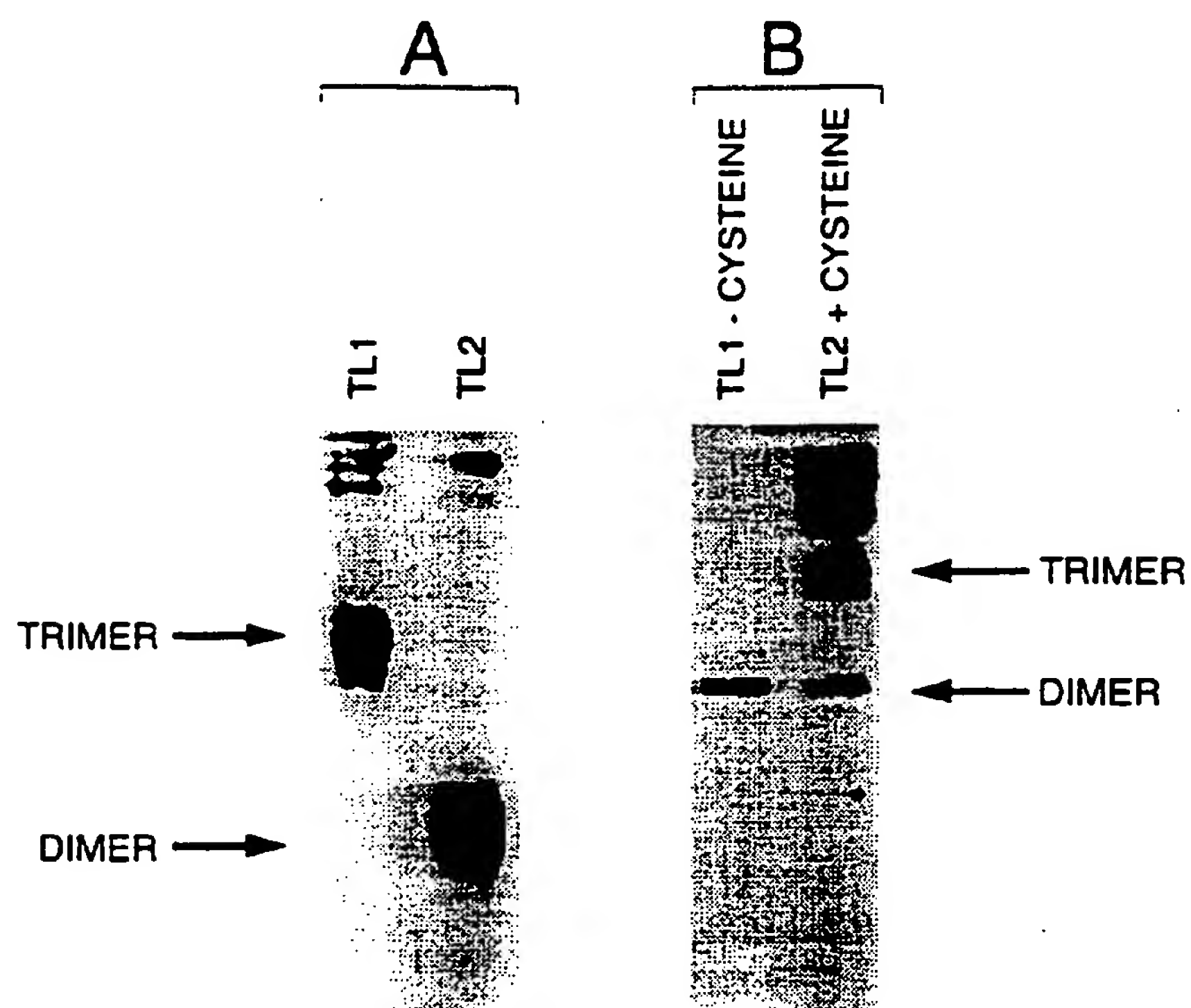
Fig.17. ENGINEERING OF TIE2 "LIGAND-BODIES"



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Fig.18.

COVALENT MULTIMERIC STRUCTURE OF
TL1 AND TL2 AND THEIR INTERCONVERSION
BY THE MUTATION OF ONE CYSTEINE



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Fig.19. TIE2-IgG binding to
immobilized TL1

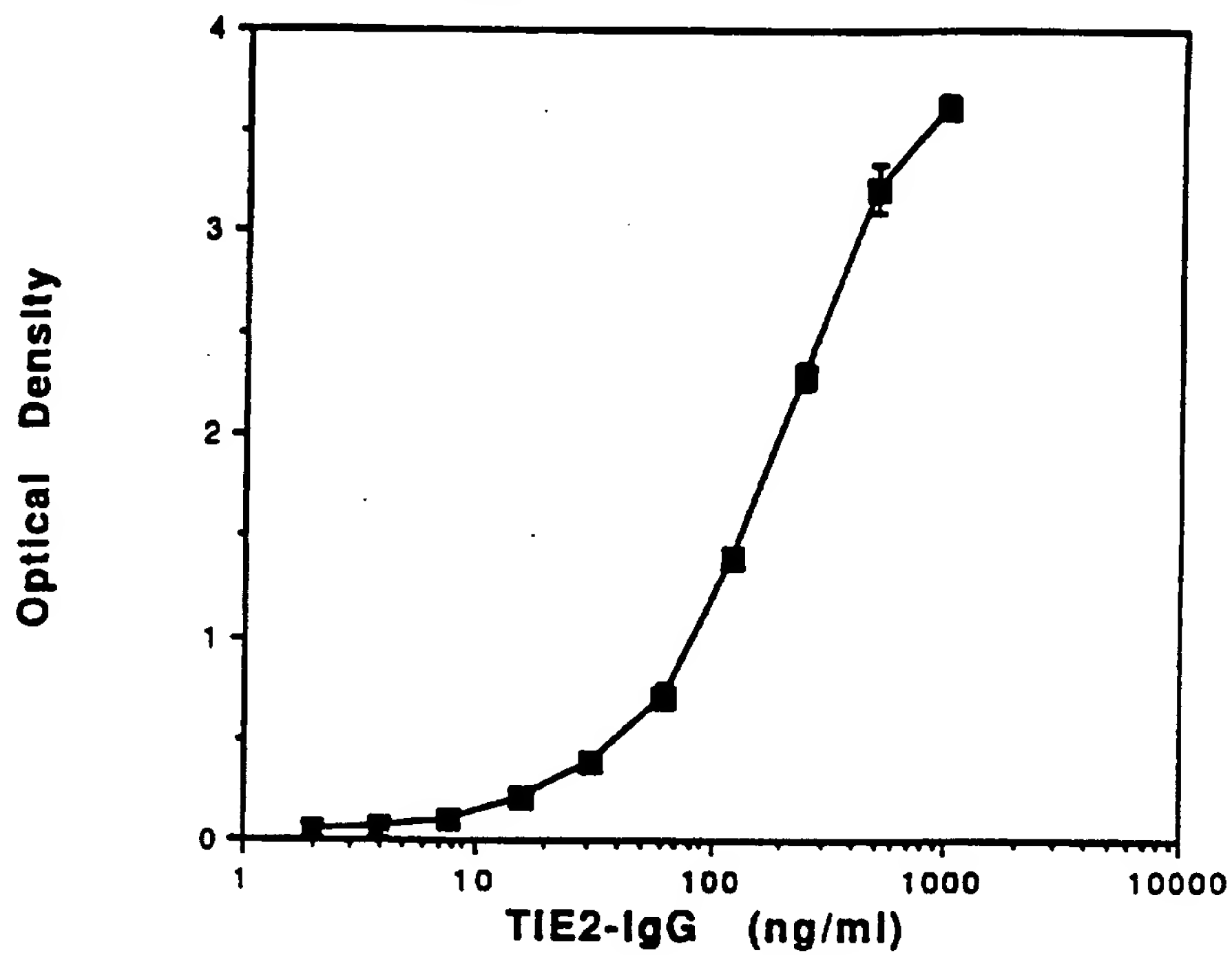
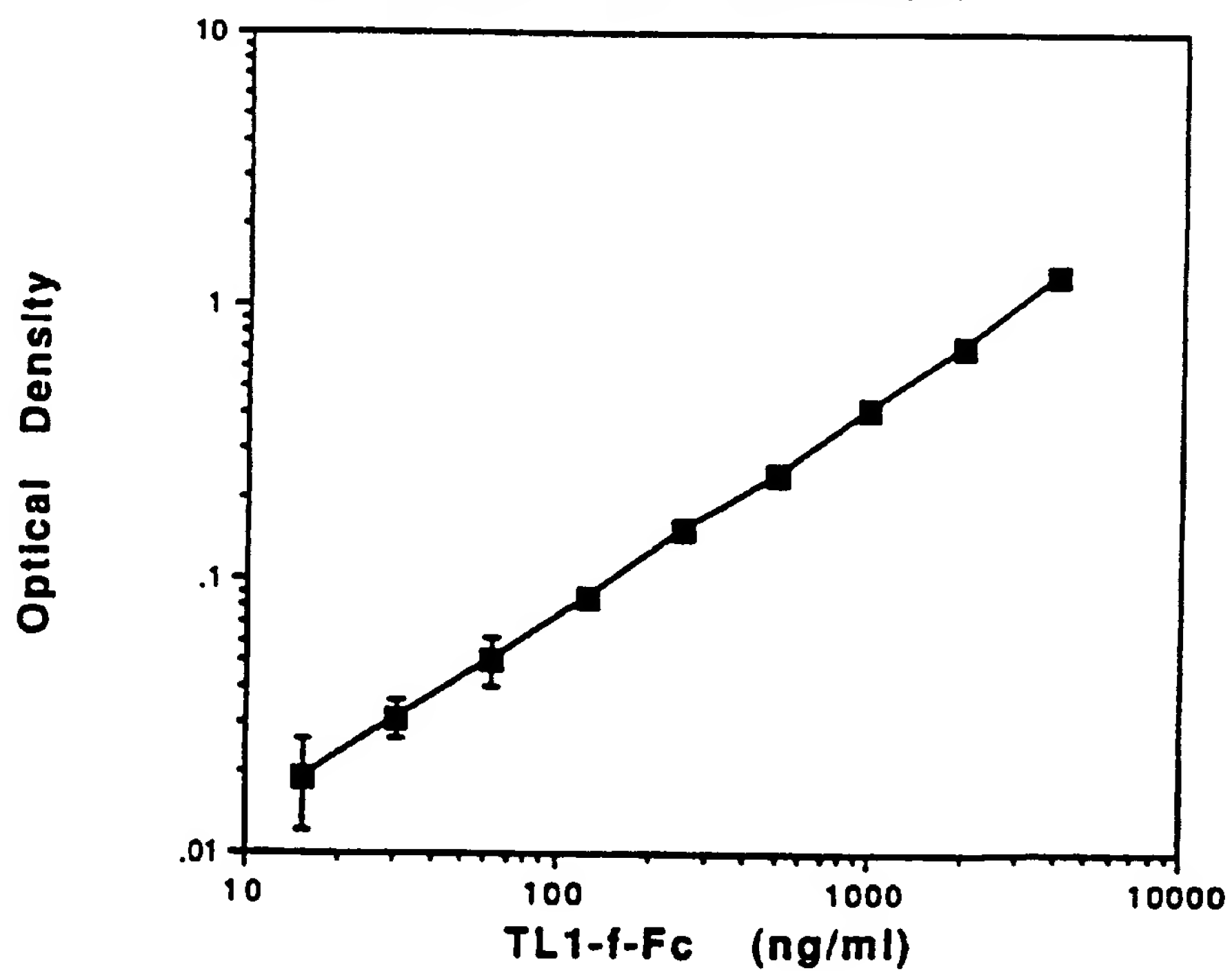


Fig.20. TL1-f-Fc binding to
immobilized Tie2 ectodomain



Applicant's or agent's file reference number REG 10-G -PCT

International application No.

INDICATIONS RELATING TO A DEPOSITED MICROORGANISM

(PCT Rule 13bis)

A. The indications made below relate to the microorganism referred to in the description on page 73, line 15 through page 74, line 2

B. IDENTIFICATION OF DEPOSIT

Further deposits are identified on an additional sheet ☒

Name of depositary institution American Type Culture Collection

Address of depositary institution (including postal code and country)
12301 Parklawn Drive
Rockville, Maryland 20852
U.S.A.

Date of deposit
October 7, 1994

Accession Number
75910

C. ADDITIONAL INDICATIONS (leave blank if not applicable) This information is continued on an additional sheet ☐

Applicant wishes that, until publication of the mention of the grant of a European patent or until the date on which the application is refused or withdrawn or is deemed to be withdrawn, the deposit shall be made available as provided in Rule 28(3) of the Implementing Regulations under the European Patent Convention only by the issue of a sample to an expert nominated by the requester (Rule 28(4) of the implementing regulations).

D. DESIGNATED STATES FOR WHICH INDICATIONS ARE MADE (if the indications are not for all designated States)

E. SEPARATE FURNISHING OF INDICATIONS (leave blank if not applicable)

The indications listed below will be submitted to the International Bureau later (specify the general nature of the indications e.g., "Accession Number of Deposit")

For receiving Office use only

☐ This sheet was received with the international application

Authorized officer

For International Bureau use only

☐ This sheet was received by the International Bureau on:

Authorized officer

Att. Dkt. No. - REG 330-G PCT

Internat'l Applic. No.:

Internat'l Filing Date: April 5, 1996

Title: TIE-2 LIGANDS, METHODS OF MAKING AND USES
THEREOF

SUPPLEMENTAL SHEET TO BOX B OF FORM PCT/RO/134

Identification of Further Deposits - In addition to the deposit indicated on the attached Form PCT/RO/134, applicant identifies the following deposits made with the American Type Culture Collection, 12301 Parklawn Drive, Rockville, MD 20852, U.S.A. and requests that they also be made available only by the issue of a sample to an expert nominated by the requester as indicated on the attached form:

Date of deposit

Accession Number

October 7, 1994

VR2484

October 26, 1994

75928

December 9, 1994

75963

INTERNATIONAL SEARCH REPORT

International application No.
PCT/US96/04806

A. CLASSIFICATION OF SUBJECT MATTER

IPC(6) :C12N 15/00, 1/20; C07K 14/52, 19/00; A61K 38/19

US CL :Please See Extra Sheet.

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

U.S. : 435/ 69.5, 320.1, 240.2, 252.3; 424/85.1; 514/2, 12; 530/351, 402; 536/23.1, 23.5

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)

APS, CAS ONLINE, MEDLINE, BIOSIS

search terms:TIE-2 ligand or protein; nucleic acid or DNA, production or isolation, treatment or administration or therapy

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A	ONCOGENE, Volume 9, Number 11, issued November 1994, Fletcher et al, "LERK-2, a Binding Protein for the Receptor-Tyrosine Kinase ELK, is Evolutionarily Conserved and Expressed in a Developmentally Regulated Pattern", pages 3241-3247, see pages 3242-3243.	1-7, 12-15, 34, 35, 39-41, 61 and 63
A	ONCOGENE, Volume 10, Number 2, issued January 1995, Kozlosky et al, "Ligands for the Receptor Tyrosine Kinases Hek and Elk: Isolation of cDNAs Encoding a Family of Proteins", pages 299-306, see entire document.	1-7, 12-15, 34, 35, 39-41, 61 and 63

☒ Further documents are listed in the continuation of Box C. ☐ See patent family annex.

A	document defining the general state of the art which is not considered to be of particular relevance	*T*	later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention
E	earlier document published on or after the international filing date	*X*	document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone
L	document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)	*Y*	document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art
O	document referring to an oral disclosure, use, exhibition or other means	*Z*	document member of the same patent family
P	document published prior to the international filing date but later than the priority date claimed		

Date of the actual completion of the international search

28 MAY 1996

Date of mailing of the international search report

08 AUG 1996

Name and mailing address of the ISA/US
Commissioner of Patents and Trademarks
Box PCT
Washington, D.C. 20231

Facsimile No. (703) 305-3230

Authorized officer

PREMA MERTZ

Telephone No. (703) 308-0196

INTERNATIONAL SEARCH REPORT

International application No.
PCT/US96/04806

C (Continuation). DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A	THE EMBO JOURNAL, Volume 13, Number 16, issued 1994, Beckmann et al, "Molecular Characterization of a Family of Ligands for Eph-Related Tyrosine Kinase Receptors", pages 3757-3762, see entire document.	1-7, 12-15, 34, 35, 39-41, 61 and 63
A	NATURE, Volume 368, issued 07 April 1994, Bartley et al, "B61 is a Ligand for the ECK Receptor Protein-Tyrosine Kinase", pages 558-560, see entire document.	1-7, 12-15, 34, 35, 39-41, 61 and 63
A	NATURE, Volume 347, issued 25 October 1990, Cattaneo et al, "Proliferation and Differentiation of Neuronal Stem Cells Regulated by Nerve Growth Factor", pages 762-765, see entire document.	1-7, 12-15, 34, 35, 39-41, 61 and 63

INTERNATIONAL SEARCH REPORT

International application No.
PCT/US96/04806

Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)

This international report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. ☐ Claims Nos.:
because they relate to subject matter not required to be searched by this Authority, namely:
2. ☐ Claims Nos.:
because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:
3. ☒ Claims Nos.: 8-11, 16-33, 36-38, 43-47, 50-60, 62, 64-69, 75
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

Please See Extra Sheet.

1. ☐ As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims.
2. ☐ As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3. ☒ As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:
1-7, 12-15, 34, 35, 39-41, 61 and 63
4. ☐ No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

Remark on Protest

☐

The additional search fees were accompanied by the applicant's protest.

☐

No protest accompanied the payment of additional search fees.

INTERNATIONAL SEARCH REPORT

International application No.

PCT/US96/04806

A. CLASSIFICATION OF SUBJECT MATTER:

US CL :

435/69.5, 320.1, 240.2, 252.3; 424/85.1; 514/2, 12; 530/351, 402; 536/23.1, 23.5

BOX II. OBSERVATIONS WHERE UNITY OF INVENTION WAS LACKING

This ISA found multiple inventions as follows:

This application contains the following inventions or groups of inventions which are not so linked as to form a single inventive concept under PCT Rule 13.1. In order for all inventions to be examined, the appropriate additional examination fees must be paid.

Group I, claims 1-3, 6-7, 12-15, 34, 35, 39, 40, and 63, drawn to a nucleic acid molecule encoding a TIE-2 ligand, a vector, a TIE-2 ligand, a host-vector system, a method of producing a TIE-2 ligand, a conjugated TIE-2 ligand, a pharmaceutical composition thereof, and a method of treating a human or animal subject by administering a TIE-2 ligand.

Group II, claims 4-5, 15, 41, and 61, drawn to a nucleic acid encoding a TIE-2 ligand antagonist and a TIE-2 ligand antagonist.

Group III, claim 42, drawn to an antibody which binds TIE-2 ligand, a pharmaceutical composition thereof, and a method of treating a human or animal subject by administering a antibody.

Group IV, claims 48-49, drawn to a method for maintaining a TIE-2 receptor expressing cell in culture.

Group V, claims 70-74, drawn to a method of treating immune system disorders(leukopenia, thrombocytopenia, anemia, enhancing bone marrow engraftment, bone marrow aplasia and a proliferative disorder of a blood forming organ) with TIE-2 ligand.

The inventions listed as Groups I-V do not relate to a single inventive concept under PCT Rule 13.1 because, under PCT Rule 13.2, they lack the same or corresponding special technical features for the following reasons:

The products/processes of Groups I-III are separate and distinct inventions in that the nucleic acid molecule encoding a TIE-2 ligand, a vector, a TIE-2 ligand, a host-vector system, a method of producing a TIE-2 ligand, a conjugated TIE-2 ligand, a pharmaceutical composition thereof, and a method of treating a human or animal subject by administering a TIE-2 ligand of Group I, the nucleic acid encoding a TIE-2 ligand antagonist and a TIE-2 ligand antagonist of Group II, the antibody which binds TIE-2 ligand, a pharmaceutical composition thereof, and a method of treating a human or animal subject by administering a antibody of Group III do not require each other for their practice and have separate functions all of which constitute the special technical features which define the contribution of each invention. Since these special technical features are not shared by each product/process and since the common features do not establish an advance over the prior art, the inventions of Groups I-IV do not form a single inventive concept within the meaning of Rule 13.2.

The processes of Groups IV-V are separate and distinct inventions in that a method for maintaining a TIE-2 receptor expressing cell in culture of Group IV, and a proliferative disorder of a blood forming organ) with TIE-2 ligand of Group V do not require each other for their practice, have separate uses and different method steps all of which constitute the special technical features which define the contribution of each invention. Since these special technical features are not shared by each process and since the common features do not establish an advance over the prior art, the inventions of Groups IV-V do not share a technical relationship and do not form a single inventive concept within the meaning of Rule 13.2.

The claims of Groups II-III are not so linked to the claims of Groups IV-V because the process steps of the methods in each of Group II-III are materially different from the process steps of each of Groups IV-V and the process steps are the special technical feature which distinguish each of the inventions from each other because they do not share a special technical feature within the meaning of Rule 13.2, so as to form a single inventive concept.

The claims of Group I are not so linked to the claims of Groups IV-V because the process steps of the methods in Group I are materially different from the process steps of each of Groups IV-V and the process steps are the special technical feature which distinguish each of the inventions from each other because they do not share a special technical feature within the meaning of Rule 13.2, so as to form a single inventive concept.